



8

SEQUENCE LISTING

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<120> Secreted and Transmembrane Polypeptides and Nucleic
Acids Encoding the Same

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<141> 2001-07-12

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<151> 1999-07-07

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<212> PRT
<213> Homo sapiens

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Arg Cys Arg Gly Leu Val Asp Lys Phe Asn Gln Gly Met Val Asp Thr
      35              40              45

Ala Lys Lys Asn Phe Gly Gly Gly Asn Thr Ala Trp Glu Glu Lys Thr
      50              55              60

Leu Ser Lys Tyr Glu Ser Ser Glu Ile Arg Leu Leu Glu Ile Leu Glu
      65              70              75              80

Gly Leu Cys Glu Ser Ser Asp Phe Glu Cys Asn Gln Met Leu Glu Ala
      85              90              95

Gln Glu Glu His Leu Glu Ala Trp Trp Leu Gln Leu Lys Ser Glu Tyr
      100             105             110

Pro Asp Leu Phe Glu Trp Phe Cys Val Lys Thr Leu Lys Val Cys Cys
      115             120             125

Ser Pro Gly Thr Tyr Gly Pro Asp Cys Leu Ala Cys Gln Gly Gly Ser
      130             135             140

Gln Arg Pro Cys Ser Gly Asn Gly His Cys Ser Gly Asp Gly Ser Arg
      145             150             155             160

Gln Gly Asp Gly Ser Cys Arg Cys His Met Gly Tyr Gln Gly Pro Leu
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Cys Thr Asp Cys Met Asp Gly Tyr Phe Ser Ser Leu Arg Asn Glu Thr
      180             185             190

His Ser Ile Cys Thr Ala Cys Asp Glu Ser Cys Lys Thr Cys Ser Gly

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| Glu Gly Ala Cys Val Asp Val Asp Glu Cys Ala Ala Glu Pro Pro Pro | | |
| 225 | 230 | 235 240 |
| Cys Ser Ala Ala Gln Phe Cys Lys Asn Ala Asn Gly Ser Tyr Thr Cys | | |
| | 245 | 250 255 |
| Glu Glu Cys Asp Ser Ser Cys Val Gly Cys Thr Gly Glu Gly Pro Gly | | |
| | 260 | 265 270 |
| Asn Cys Lys Glu Cys Ile Ser Gly Tyr Ala Arg Glu His Gly Gln Cys | | |
| | 275 | 280 285 |
| Ala Asp Val Asp Glu Cys Ser Leu Ala Glu Lys Thr Cys Val Arg Lys | | |
| | 290 | 295 300 |
| Asn Glu Asn Cys Tyr Asn Thr Pro Gly Ser Tyr Val Cys Val Cys Pro | | |
| 305 | 310 | 315 320 |
| Asp Gly Phe Glu Glu Thr Glu Asp Ala Cys Val Pro Pro Ala Glu Ala | | |
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| Glu Ala Thr Glu Gly Glu Ser Pro Thr Gln Leu Pro Ser Arg Glu Asp | | |
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<210> 3
 <211> 2206
 <212> DNA
 <213> Homo sapiens

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| aacagccctg | gctgagggag | ctgcagcgca | gcagagtatc | tgacggcgcc | aggttgcgta | 180 |
| ggtgcggcac | gaggagtttt | cccggcagcg | aggaggtcct | gagcagcatg | gcccggagga | 240 |
| gcgccttccc | tgccgcccg | ctctggctct | ggagcatcct | cctgtgcctg | ctggcactgc | 300 |
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| caagagtact | cataggattt | gaagaagata | tcctgattgt | ttcagagggg | aaaatggcac | 420 |
| cttttacaca | tgatttcaga | aaagcgcaac | agagaatgcc | agctattcct | gtcaatatcc | 480 |
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| ccttgcgctc | cctggataaa | ggcatcatgg | cagatccaac | cgtcaatgtc | cctctgctgg | 600 |
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| tgactcctgg | tttctgcatc | tgcccacctg | gattctatgg | agtgaactgt | gacaaagcaa | 960 |
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 <212> PRT
 <213> Homo sapiens

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      20              25              30

Glu Glu Ser Leu Tyr Leu Trp Ile Asp Ala His Gln Ala Arg Val Leu
      35              40              45

Ile Gly Phe Glu Glu Asp Ile Leu Ile Val Ser Glu Gly Lys Met Ala
      50              55              60

Pro Phe Thr His Asp Phe Arg Lys Ala Gln Gln Arg Met Pro Ala Ile
      65              70              75              80

Pro Val Asn Ile His Ser Met Asn Phe Thr Trp Gln Ala Ala Gly Gln
      85              90              95

Ala Glu Tyr Phe Tyr Glu Phe Leu Ser Leu Arg Ser Leu Asp Lys Gly
      100              105              110

Ile Met Ala Asp Pro Thr Val Asn Val Pro Leu Leu Gly Thr Val Pro
      115              120              125

His Lys Ala Ser Val Val Gln Val Gly Phe Pro Cys Leu Gly Lys Gln
      130              135              140

Asp Gly Val Ala Ala Phe Glu Val Asp Val Ile Val Met Asn Ser Glu
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Gly Asn Thr Ile Leu Gln Thr Pro Gln Asn Ala Ile Phe Phe Lys Thr
      165              170              175

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Cys Gln Gln Ala Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys
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 Asn Glu Arg Arg Ile Cys Glu Cys Pro Asp Gly Phe His Gly Pro His
 195 200 205
 Cys Glu Lys Ala Leu Cys Thr Pro Arg Cys Met Asn Gly Gly Leu Cys
 210 215 220
 Val Thr Pro Gly Phe Cys Ile Cys Pro Pro Gly Phe Tyr Gly Val Asn
 225 230 235 240
 Cys Asp Lys Ala Asn Cys Ser Thr Thr Cys Phe Asn Gly Gly Thr Cys
 245 250 255
 Phe Tyr Pro Gly Lys Cys Ile Cys Pro Pro Gly Leu Glu Gly Glu Gln
 260 265 270
 Cys Glu Ile Ser Lys Cys Pro Gln Pro Cys Arg Asn Gly Gly Lys Cys
 275 280 285
 Ile Gly Lys Ser Lys Cys Lys Cys Ser Lys Gly Tyr Gln Gly Asp Leu
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 Cys Ser Lys Pro Val Cys Glu Pro Gly Cys Gly Ala His Gly Thr Cys
 305 310 315 320
 His Glu Pro Asn Lys Cys Gln Cys Gln Glu Gly Trp His Gly Arg His
 325 330 335
 Cys Asn Lys Arg Tyr Glu Ala Ser Leu Ile His Ala Leu Arg Pro Ala
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<210> 5

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

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<210> 6

<211> 21

<212> DNA

<213> Artificial Sequence

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 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

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<210> 7
 <211> 22
 <212> DNA
 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 7
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<210> 8
 <211> 49
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
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<210> 9
 <211> 22
 <212> DNA
 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Synthetic
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<400> 9
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<210> 10
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<220>
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<400> 10
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 <211> 2197

<212> DNA
<213> Homo sapiens

<400> 11

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<210> 12
<211> 164
<212> PRT
<213> Homo sapiens

<400> 12

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      20                25                30

Pro Gly Leu His Leu Arg Gly Ile Arg Asp Ala Gly Gly Arg Tyr Cys
      35                40                45
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Gln Glu Gln Asp Leu Cys Cys Arg Gly Arg Ala Asp Asp Cys Ala Leu
 50 55 60
 Pro Tyr Leu Gly Ala Ile Cys Tyr Cys Asp Leu Phe Cys Asn Arg Thr
 65 70 75 80
 Val Ser Asp Cys Cys Pro Asp Phe Trp Asp Phe Cys Leu Gly Val Pro
 85 90 95
 Pro Pro Phe Pro Pro Ile Gln Gly Cys Met His Gly Gly Arg Ile Tyr
 100 105 110
 Pro Val Leu Gly Thr Tyr Trp Asp Asn Cys Asn Arg Cys Thr Cys Gln
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 Glu Asn Arg Gln Trp His Gly Gly Ser Arg His Asp Gln Ser His Gln
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 Pro Gly Gln Leu Trp Leu Ala Gly Trp Glu Pro Gln Arg Leu Leu Gly
 145 150 155 160
 His Asp Pro Gly

<210> 13
 <211> 533
 <212> DNA
 <213> Homo sapiens

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<223> a, t, c or g

<400> 13

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<210> 14

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<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

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<210> 15

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<212> DNA

<213> Artificial Sequence

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oligonucleotide probe

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<210> 16

<211> 50

<212> DNA

<213> Artificial Sequence

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oligonucleotide probe

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<210> 17

<211> 960

<212> DNA

<213> Homo sapiens

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<210> 18
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 <213> Homo sapiens

<400> 18

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| Met | Thr | His | Arg | Thr | Thr | Thr | Trp | Ala | Arg | Arg | Thr | Ser | Arg | Ala | Val |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| | | | | | | | | | | | | | | | |
| Thr | Pro | Thr | Cys | Ala | Thr | Pro | Ala | Gly | Pro | Met | Pro | Cys | Ser | Arg | Leu |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| | | | | | | | | | | | | | | | |
| Pro | Pro | Ser | Leu | Arg | Cys | Ser | Leu | His | Ser | Ala | Cys | Cys | Ser | Gly | Asp |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| | | | | | | | | | | | | | | | |
| Pro | Ala | Ser | Tyr | Arg | Leu | Trp | Gly | Ala | Pro | Leu | Gln | Pro | Thr | Leu | Gly |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| | | | | | | | | | | | | | | | |
| Val | Val | Pro | Gln | Ala | Ser | Val | Pro | Leu | Leu | Thr | Asp | Leu | Ala | Gln | Trp |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| | | | | | | | | | | | | | | | |
| Glu | Pro | Val | Leu | Val | Pro | Glu | Ala | His | Pro | Asn | Ala | Ser | Leu | Thr | Met |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| | | | | | | | | | | | | | | | |
| Tyr | Val | Cys | Thr | Pro | Val | Pro | His | Pro | Asp | Pro | Pro | Met | Ala | Leu | Ser |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| | | | | | | | | | | | | | | | |
| Arg | Thr | Pro | Thr | Arg | Gln | Ile | Ser | Ser | Ser | Asp | Thr | Asp | Pro | Pro | Ala |
| | | 115 | | | | 120 | | | | | | 125 | | | |
| | | | | | | | | | | | | | | | |
| Asp | Gly | Pro | Ser | Asn | Pro | Leu | Cys | Cys | Cys | Phe | His | Gly | Pro | Ala | Phe |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| | | | | | | | | | | | | | | | |
| Ser | Thr | Leu | Asn | Pro | Val | Leu | Arg | His | Leu | Phe | Pro | Gln | Glu | Ala | Phe |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| | | | | | | | | | | | | | | | |
| Pro | Ala | His | Pro | Ile | Tyr | Asp | Leu | Ser | Gln | Val | Trp | Ser | Val | Val | Ser |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| | | | | | | | | | | | | | | | |
| Pro | Ala | Pro | Ser | Arg | Gly | Gln | Ala | Leu | Arg | Arg | Ala | Gln | | | |
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<210> 19
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 19
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<210> 20
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 20
tgcacaagtc ggtgtcacag cacg 24

<210> 21
<211> 44
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 21
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<210> 22
<211> 1200
<212> DNA
<213> Homo sapiens

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<210> 23
 <211> 205
 <212> PRT
 <213> Homo sapiens

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 35 40 45
 Arg Thr Ser Gly Lys His Val Gln Val Thr Gly Arg Arg Ile Ser Ala
 50 55 60
 Thr Ala Glu Asp Gly Asn Lys Phe Ala Lys Leu Ile Val Glu Thr Asp
 65 70 75 80
 Thr Phe Gly Ser Arg Val Arg Ile Lys Gly Ala Glu Ser Glu Lys Tyr
 85 90 95
 Ile Cys Met Asn Lys Arg Gly Lys Leu Ile Gly Lys Pro Ser Gly Lys
 100 105 110
 Ser Lys Asp Cys Val Phe Thr Glu Ile Val Leu Glu Asn Asn Tyr Thr
 115 120 125
 Ala Phe Gln Asn Ala Arg His Glu Gly Trp Phe Met Ala Phe Thr Arg
 130 135 140
 Gln Gly Arg Pro Arg Gln Ala Ser Arg Ser Arg Gln Asn Gln Arg Glu
 145 150 155 160
 Ala His Phe Ile Lys Arg Leu Tyr Gln Gly Gln Leu Pro Phe Pro Asn
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 195 200 205

<210> 24
 <211> 28
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 24
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<210> 25
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 25
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<210> 26
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<220>
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<222> (21)..(21)
<223> a, t, c or g

<400> 26
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<210> 27
<211> 2479
<212> DNA
<213> Homo sapiens

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2479

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<210> 28
 <211> 660
 <212> PRT
 <213> Homo sapiens

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<400> 28
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Lys Ser Trp Leu Ile Ile Ser Leu Gly Leu Tyr Ser Gln Val Ser Lys
      20                      25          30

Leu Leu Ala Cys Pro Ser Val Cys Arg Cys Asp Arg Asn Phe Val Tyr
      35                      40          45

Cys Asn Glu Arg Ser Leu Thr Ser Val Pro Leu Gly Ile Pro Glu Gly
      50                      55          60

Val Thr Val Leu Tyr Leu His Asn Asn Gln Ile Asn Asn Ala Gly Phe
      65                      70          75          80

Pro Ala Glu Leu His Asn Val Gln Ser Val His Thr Val Tyr Leu Tyr
      85                      90          95

Gly Asn Gln Leu Asp Glu Phe Pro Met Asn Leu Pro Lys Asn Val Arg
      100                     105          110

Val Leu His Leu Gln Glu Asn Asn Ile Gln Thr Ile Ser Arg Ala Ala
      115                     120          125

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| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Ala | Gln | Leu | Leu | Lys | Leu | Glu | Glu | Leu | His | Leu | Asp | Asp | Asn | Ser | 130 | 135 | 140 |
| Ile | Ser | Thr | Val | Gly | Val | Glu | Asp | Gly | Ala | Phe | Arg | Glu | Ala | Ile | Ser | 145 | 150 | 155 |
| Leu | Lys | Leu | Leu | Phe | Leu | Ser | Lys | Asn | His | Leu | Ser | Ser | Val | Pro | Val | 165 | 170 | 175 |
| Gly | Leu | Pro | Val | Asp | Leu | Gln | Glu | Leu | Arg | Val | Asp | Glu | Asn | Arg | Ile | 180 | 185 | 190 |
| Ala | Val | Ile | Ser | Asp | Met | Ala | Phe | Gln | Asn | Leu | Thr | Ser | Leu | Glu | Arg | 195 | 200 | 205 |
| Leu | Ile | Val | Asp | Gly | Asn | Leu | Leu | Thr | Asn | Lys | Gly | Ile | Ala | Glu | Gly | 210 | 215 | 220 |
| Thr | Phe | Ser | His | Leu | Thr | Lys | Leu | Lys | Glu | Phe | Ser | Ile | Val | Arg | Asn | 225 | 230 | 235 |
| Ser | Leu | Ser | His | Pro | Pro | Pro | Asp | Leu | Pro | Gly | Thr | His | Leu | Ile | Arg | 245 | 250 | 255 |
| Leu | Tyr | Leu | Gln | Asp | Asn | Gln | Ile | Asn | His | Ile | Pro | Leu | Thr | Ala | Phe | 260 | 265 | 270 |
| Ser | Asn | Leu | Arg | Lys | Leu | Glu | Arg | Leu | Asp | Ile | Ser | Asn | Asn | Gln | Leu | 275 | 280 | 285 |
| Arg | Met | Leu | Thr | Gln | Gly | Val | Phe | Asp | Asn | Leu | Ser | Asn | Leu | Lys | Gln | 290 | 295 | 300 |
| Leu | Thr | Ala | Arg | Asn | Asn | Pro | Trp | Phe | Cys | Asp | Cys | Ser | Ile | Lys | Trp | 305 | 310 | 315 |
| Val | Thr | Glu | Trp | Leu | Lys | Tyr | Ile | Pro | Ser | Ser | Leu | Asn | Val | Arg | Gly | 325 | 330 | 335 |
| Phe | Met | Cys | Gln | Gly | Pro | Glu | Gln | Val | Arg | Gly | Met | Ala | Val | Arg | Glu | 340 | 345 | 350 |
| Leu | Asn | Met | Asn | Leu | Leu | Ser | Cys | Pro | Thr | Thr | Thr | Pro | Gly | Leu | Pro | 355 | 360 | 365 |
| Leu | Phe | Thr | Pro | Ala | Pro | Ser | Thr | Ala | Ser | Pro | Thr | Thr | Gln | Pro | Pro | 370 | 375 | 380 |
| Thr | Leu | Ser | Ile | Pro | Asn | Pro | Ser | Arg | Ser | Tyr | Thr | Pro | Pro | Thr | Pro | 385 | 390 | 395 |
| Thr | Thr | Ser | Lys | Leu | Pro | Thr | Ile | Pro | Asp | Trp | Asp | Gly | Arg | Glu | Arg | 405 | 410 | 415 |
| Val | Thr | Pro | Pro | Ile | Ser | Glu | Arg | Ile | Gln | Leu | Ser | Ile | His | Phe | Val | 420 | 425 | 430 |

Asn Asp Thr Ser Ile Gln Val Ser Trp Leu Ser Leu Phe Thr Val Met
 435 440 445
 Ala Tyr Lys Leu Thr Trp Val Lys Met Gly His Ser Leu Val Gly Gly
 450 455 460
 Ile Val Gln Glu Arg Ile Val Ser Gly Glu Lys Gln His Leu Ser Leu
 465 470 475 480
 Val Asn Leu Glu Pro Arg Ser Thr Tyr Arg Ile Cys Leu Val Pro Leu
 485 490 495
 Asp Ala Phe Asn Tyr Arg Ala Val Glu Asp Thr Ile Cys Ser Glu Ala
 500 505 510
 Thr Thr His Ala Ser Tyr Leu Asn Asn Gly Ser Asn Thr Ala Ser Ser
 515 520 525
 His Glu Gln Thr Thr Ser His Ser Met Gly Ser Pro Phe Leu Leu Ala
 530 535 540
 Gly Leu Ile Gly Gly Ala Val Ile Phe Val Leu Val Val Leu Leu Ser
 545 550 555 560
 Val Phe Cys Trp His Met His Lys Lys Gly Arg Tyr Thr Ser Gln Lys
 565 570 575
 Trp Lys Tyr Asn Arg Gly Arg Arg Lys Asp Asp Tyr Cys Glu Ala Gly
 580 585 590
 Thr Lys Lys Asp Asn Ser Ile Leu Glu Met Thr Glu Thr Ser Phe Gln
 595 600 605
 Ile Val Ser Leu Asn Asn Asp Gln Leu Leu Lys Gly Asp Phe Arg Leu
 610 615 620
 Gln Pro Ile Tyr Thr Pro Asn Gly Gly Ile Asn Tyr Thr Asp Cys His
 625 630 635 640
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 His Cys His Thr
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<210> 29

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 29

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<210> 30
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 30
gcaggacaac cagataaacc ac 22

<210> 31
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
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oligonucleotide probe

<400> 31
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<210> 32
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<212> DNA
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<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

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<210> 33
<211> 3449
<212> DNA
<213> Homo sapiens

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 <212> PRT
 <213> Homo sapiens

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 20 25 30
 Ser Arg Gly Arg His Ala Arg Thr His Pro Gln Thr Ala Leu Leu Glu

| 35 | | | | | 40 | | | | | 45 | | | | | |
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| Ser | Ser | Cys | Glu | Asn | Lys | Arg | Ala | Asp | Leu | Val | Phe | Ile | Ile | Asp | Ser |
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| Ser | Arg | Ser | Val | Asn | Thr | His | Asp | Tyr | Ala | Lys | Val | Lys | Glu | Phe | Ile |
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| Val | Asp | Ile | Leu | Gln | Phe | Leu | Asp | Ile | Gly | Pro | Asp | Val | Thr | Arg | Val |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| Gly | Leu | Leu | Gln | Tyr | Gly | Ser | Thr | Val | Lys | Asn | Glu | Phe | Ser | Leu | Lys |
| | | | 100 | | | | | 105 | | | | | 110 | | |
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| | | 115 | | | | | 120 | | | | | 125 | | | |
| His | Leu | Ser | Thr | Gly | Thr | Met | Thr | Gly | Leu | Ala | Ile | Gln | Tyr | Ala | Leu |
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| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Val | Pro | Arg | Val | Ile | Met | Ile | Val | Thr | Asp | Gly | Arg | Pro | Gln | Asp | Ser |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Val | Ala | Glu | Val | Ala | Ala | Lys | Ala | Arg | Asp | Thr | Gly | Ile | Leu | Ile | Phe |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Ala | Ile | Gly | Val | Gly | Gln | Val | Asp | Phe | Asn | Thr | Leu | Lys | Ser | Ile | Gly |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Ser | Glu | Pro | His | Glu | Asp | His | Val | Phe | Leu | Val | Ala | Asn | Phe | Ser | Gln |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Ile | Glu | Thr | Leu | Thr | Ser | Val | Phe | Gln | Lys | Lys | Leu | Cys | Thr | Ala | His |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Met | Cys | Ser | Thr | Leu | Glu | His | Asn | Cys | Ala | His | Phe | Cys | Ile | Asn | Ile |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Pro | Gly | Ser | Tyr | Val | Cys | Arg | Cys | Lys | Gln | Gly | Tyr | Ile | Leu | Asn | Ser |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Asp | Gln | Thr | Thr | Cys | Arg | Ile | Gln | Asp | Leu | Cys | Ala | Met | Glu | Asp | His |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Asn | Cys | Glu | Gln | Leu | Cys | Val | Asn | Val | Pro | Gly | Ser | Phe | Val | Cys | Gln |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Cys | Tyr | Ser | Gly | Tyr | Ala | Leu | Ala | Glu | Asp | Gly | Lys | Arg | Cys | Val | Ala |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Val | Asp | Tyr | Cys | Ala | Ser | Glu | Asn | His | Gly | Cys | Glu | His | Glu | Cys | Val |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Asn | Ala | Asp | Gly | Ser | Tyr | Leu | Cys | Gln | Cys | His | Glu | Gly | Phe | Ala | Leu |

| 340 | | | | | 345 | | | | | 350 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Pro | Asp | Glu | Lys | Thr | Cys | Thr | Arg | Ile | Asn | Tyr | Cys | Ala | Leu | Asn |
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| Lys | Pro | Gly | Cys | Glu | His | Glu | Cys | Val | Asn | Met | Glu | Glu | Ser | Tyr | Tyr |
| | 370 | | | | | 375 | | | | | 380 | | | | |
| Cys | Arg | Cys | His | Arg | Gly | Tyr | Thr | Leu | Asp | Pro | Asn | Gly | Lys | Thr | Cys |
| 385 | | | | | | 390 | | | | | 395 | | | | 400 |
| Ser | Arg | Val | Asp | His | Cys | Ala | Gln | Gln | Asp | His | Gly | Cys | Glu | Gln | Leu |
| | | | | 405 | | | | | 410 | | | | | 415 | |
| Cys | Leu | Asn | Thr | Glu | Asp | Ser | Phe | Val | Cys | Gln | Cys | Ser | Glu | Gly | Phe |
| | | | 420 | | | | | 425 | | | | | 430 | | |
| Leu | Ile | Asn | Glu | Asp | Leu | Lys | Thr | Cys | Ser | Arg | Val | Asp | Tyr | Cys | Leu |
| | | 435 | | | | | 440 | | | | | 445 | | | |
| Leu | Ser | Asp | His | Gly | Cys | Glu | Tyr | Ser | Cys | Val | Asn | Met | Asp | Arg | Ser |
| | 450 | | | | | 455 | | | | | 460 | | | | |
| Phe | Ala | Cys | Gln | Cys | Pro | Glu | Gly | His | Val | Leu | Arg | Ser | Asp | Gly | Lys |
| 465 | | | | | | 470 | | | | | 475 | | | | 480 |
| Thr | Cys | Ala | Lys | Leu | Asp | Ser | Cys | Ala | Leu | Gly | Asp | His | Gly | Cys | Glu |
| | | | | 485 | | | | | 490 | | | | | 495 | |
| His | Ser | Cys | Val | Ser | Ser | Glu | Asp | Ser | Phe | Val | Cys | Gln | Cys | Phe | Glu |
| | | | 500 | | | | | 505 | | | | | 510 | | |
| Gly | Tyr | Ile | Leu | Arg | Glu | Asp | Gly | Lys | Thr | Cys | Arg | Arg | Lys | Asp | Val |
| | | 515 | | | | | 520 | | | | | 525 | | | |
| Cys | Gln | Ala | Ile | Asp | His | Gly | Cys | Glu | His | Ile | Cys | Val | Asn | Ser | Asp |
| | 530 | | | | | 535 | | | | | 540 | | | | |
| Asp | Ser | Tyr | Thr | Cys | Glu | Cys | Leu | Glu | Gly | Phe | Arg | Leu | Ala | Glu | Asp |
| 545 | | | | | | 550 | | | | | 555 | | | | 560 |
| Gly | Lys | Arg | Cys | Arg | Arg | Lys | Asp | Val | Cys | Lys | Ser | Thr | His | His | Gly |
| | | | | 565 | | | | | 570 | | | | | 575 | |
| Cys | Glu | His | Ile | Cys | Val | Asn | Asn | Gly | Asn | Ser | Tyr | Ile | Cys | Lys | Cys |
| | | | 580 | | | | | 585 | | | | | 590 | | |
| Ser | Glu | Gly | Phe | Val | Leu | Ala | Glu | Asp | Gly | Arg | Arg | Cys | Lys | Lys | Cys |
| | | 595 | | | | | 600 | | | | | 605 | | | |
| Thr | Glu | Gly | Pro | Ile | Asp | Leu | Val | Phe | Val | Ile | Asp | Gly | Ser | Lys | Ser |
| | | | | | | 615 | | | | | 620 | | | | |
| Leu | Gly | Glu | Glu | Asn | Phe | Glu | Val | Val | Lys | Gln | Phe | Val | Thr | Gly | Ile |
| 625 | | | | | | 630 | | | | | 635 | | | | 640 |
| Ile | Asp | Ser | Leu | Thr | Ile | Ser | Pro | Lys | Ala | Ala | Arg | Val | Gly | Leu | Leu |

| 645 | | | | | 650 | | | | | 655 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gln | Tyr | Ser | Thr | Gln | Val | His | Thr | Glu | Phe | Thr | Leu | Arg | Asn | Phe | Asn |
| | | | 660 | | | | | 665 | | | | | 670 | | |
| Ser | Ala | Lys | Asp | Met | Lys | Lys | Ala | Val | Ala | His | Met | Lys | Tyr | Met | Gly |
| | | 675 | | | | | 680 | | | | | 685 | | | |
| Lys | Gly | Ser | Met | Thr | Gly | Leu | Ala | Leu | Lys | His | Met | Phe | Glu | Arg | Ser |
| | 690 | | | | | 695 | | | | | 700 | | | | |
| Phe | Thr | Gln | Gly | Glu | Gly | Ala | Arg | Pro | Leu | Ser | Thr | Arg | Val | Pro | Arg |
| 705 | | | | | | 710 | | | | | 715 | | | | 720 |
| Ala | Ala | Ile | Val | Phe | Thr | Asp | Gly | Arg | Ala | Gln | Asp | Asp | Val | Ser | Glu |
| | | | | 725 | | | | | 730 | | | | | 735 | |
| Trp | Ala | Ser | Lys | Ala | Lys | Ala | Asn | Gly | Ile | Thr | Met | Tyr | Ala | Val | Gly |
| | | | 740 | | | | | 745 | | | | | 750 | | |
| Val | Gly | Lys | Ala | Ile | Glu | Glu | Glu | Leu | Gln | Glu | Ile | Ala | Ser | Glu | Pro |
| | | 755 | | | | | 760 | | | | | 765 | | | |
| Thr | Asn | Lys | His | Leu | Phe | Tyr | Ala | Glu | Asp | Phe | Ser | Thr | Met | Asp | Glu |
| | 770 | | | | | 775 | | | | | 780 | | | | |
| Ile | Ser | Glu | Lys | Leu | Lys | Lys | Gly | Ile | Cys | Glu | Ala | Leu | Glu | Asp | Ser |
| 785 | | | | 790 | | | | | | | 795 | | | | 800 |
| Asp | Gly | Arg | Gln | Asp | Ser | Pro | Ala | Gly | Glu | Leu | Pro | Lys | Thr | Val | Gln |
| | | | | 805 | | | | | 810 | | | | | 815 | |
| Gln | Pro | Thr | Glu | Ser | Glu | Pro | Val | Thr | Ile | Asn | Ile | Gln | Asp | Leu | Leu |
| | | | 820 | | | | | 825 | | | | | 830 | | |
| Ser | Cys | Ser | Asn | Phe | Ala | Val | Gln | His | Arg | Tyr | Leu | Phe | Glu | Glu | Asp |
| | | 835 | | | | | 840 | | | | | 845 | | | |
| Asn | Leu | Leu | Arg | Ser | Thr | Gln | Lys | Leu | Ser | His | Ser | Thr | Lys | Pro | Ser |
| | 850 | | | | | 855 | | | | | 860 | | | | |
| Gly | Ser | Pro | Leu | Glu | Glu | Lys | His | Asp | Gln | Cys | Lys | Cys | Glu | Asn | Leu |
| 865 | | | | 870 | | | | | | | 875 | | | | 880 |
| Ile | Met | Phe | Gln | Asn | Leu | Ala | Asn | Glu | Glu | Val | Arg | Lys | Leu | Thr | Gln |
| | | | 885 | | | | | | 890 | | | | | 895 | |
| Arg | Leu | Glu | Glu | Met | Thr | Gln | Arg | Met | Glu | Ala | Leu | Glu | Asn | Arg | Leu |
| | | | 900 | | | | | 905 | | | | | 910 | | |
| Arg | Tyr | Arg | | | | | | | | | | | | | |
| | | | 915 | | | | | | | | | | | | |

<210> 35
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 35

gtgaccctgg ttgtgaatac tcc

23

<210> 36

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 36

acagccatgg tctatagctt gg

22

<210> 37

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 37

gcctgtcagt gtccctgaggg acacgtgctc cgcagcgatg ggaag

45

<210> 38

<211> 1813

<212> DNA

<213> Homo sapiens

<400> 38

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 <213> Homo sapiens

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      20           25           30

Leu His Leu Pro Ala Asn Arg Leu Gln Ala Val Glu Gly Gly Glu Val
      35           40           45

Val Leu Pro Ala Trp Tyr Thr Leu His Gly Glu Val Ser Ser Ser Gln
      50           55           60

Pro Trp Glu Val Pro Phe Val Met Trp Phe Phe Lys Gln Lys Glu Lys
      65           70           75           80

Glu Asp Gln Val Leu Ser Tyr Ile Asn Gly Val Thr Thr Ser Lys Pro
      85           90           95

Gly Val Ser Leu Val Tyr Ser Met Pro Ser Arg Asn Leu Ser Leu Arg
      100          105          110

Leu Glu Gly Leu Gln Glu Lys Asp Ser Gly Pro Tyr Ser Cys Ser Val
      115          120          125

Asn Val Gln Asp Lys Gln Gly Lys Ser Arg Gly His Ser Ile Lys Thr
      130          135          140

Leu Glu Leu Asn Val Leu Val Pro Pro Ala Pro Pro Ser Cys Arg Leu
      145          150          155          160

Gln Gly Val Pro His Val Gly Ala Asn Val Thr Leu Ser Cys Gln Ser
      165          170          175

Pro Arg Ser Lys Pro Ala Val Gln Tyr Gln Trp Asp Arg Gln Leu Pro
      180          185          190

Ser Phe Gln Thr Phe Phe Ala Pro Ala Leu Asp Val Ile Arg Gly Ser
      195          200          205

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Leu Ser Leu Thr Asn Leu Ser Ser Ser Met Ala Gly Val Tyr Val Cys
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 Lys Ala His Asn Glu Val Gly Thr Ala Gln Cys Asn Val Thr Leu Glu
 225 230 235 240
 Val Ser Thr Gly Pro Gly Ala Ala Val Val Ala Gly Ala Val Val Gly
 245 250 255
 Thr Leu Val Gly Leu Gly Leu Leu Ala Gly Leu Val Leu Leu Tyr His
 260 265 270
 Arg Arg Gly Lys Ala Leu Glu Glu Pro Ala Asn Asp Ile Lys Glu Asp
 275 280 285
 Ala Ile Ala Pro Arg Thr Leu Pro Trp Pro Lys Ser Ser Asp Thr Ile
 290 295 300
 Ser Lys Asn Gly Thr Leu Ser Ser Val Thr Ser Ala Arg Ala Leu Arg
 305 310 315 320
 Pro Pro His Gly Pro Pro Arg Pro Gly Ala Leu Thr Pro Thr Pro Ser
 325 330 335
 Leu Ser Ser Gln Ala Leu Pro Ser Pro Arg Leu Pro Thr Thr Asp Gly
 340 345 350
 Ala His Pro Gln Pro Ile Ser Pro Ile Pro Gly Gly Val Ser Ser Ser
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 370 375 380
 Gln Ala Gly Ser Leu Val
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<210> 40

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 40

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22

<210> 41

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 41
 attgtgggcc ttgcagacat agac 24

<210> 42
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 42
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<210> 43
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 43
 gtgtgacaca gcgtgggc 18

<210> 44
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 44
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<210> 45
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 45
 cagcagcttc agccaccagg agtgg 25

<210> 46
 <211> 24
 <212> DNA
 <213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 46

ctgagccgtg ggctgcagtc tcgc

24

<210> 47

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 47

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45

<210> 48

<211> 2822

<212> DNA

<213> Homo sapiens

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agacttctgt ttgctaaatc tgtttctttt tctaataattc taaaaaaaaa aaaaagggtt 2760
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aa

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<210> 49
 <211> 690
 <212> PRT
 <213> Homo sapiens

<400> 49
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 Ser Tyr Thr Gln Asn Cys Thr Lys Thr Pro Cys Leu Pro Asn Ala Lys
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 35 40 45
 Ser Gly Asn Gly Val Thr Ile Cys Glu Asp Asp Asn Glu Cys Gly Asn
 50 55 60
 Leu Thr Gln Ser Cys Gly Glu Asn Ala Asn Cys Thr Asn Thr Glu Gly
 65 70 75 80
 Ser Tyr Tyr Cys Met Cys Val Pro Gly Phe Arg Ser Ser Ser Asn Gln
 85 90 95
 Asp Arg Phe Ile Thr Asn Asp Gly Thr Val Cys Ile Glu Asn Val Asn
 100 105 110
 Ala Asn Cys His Leu Asp Asn Val Cys Ile Ala Ala Asn Ile Asn Lys
 115 120 125
 Thr Leu Thr Lys Ile Arg Ser Ile Lys Glu Pro Val Ala Leu Leu Gln
 130 135 140
 Glu Val Tyr Arg Asn Ser Val Thr Asp Leu Ser Pro Thr Asp Ile Ile
 145 150 155 160
 Thr Tyr Ile Glu Ile Leu Ala Glu Ser Ser Ser Leu Leu Gly Tyr Lys
 165 170 175
 Asn Asn Thr Ile Ser Ala Lys Asp Thr Leu Ser Asn Ser Thr Leu Thr

| 180 | | | | | 185 | | | | | 190 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Phe | Val | Lys | Thr | Val | Asn | Asn | Phe | Val | Gln | Arg | Asp | Thr | Phe | Val |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Val | Trp | Asp | Lys | Leu | Ser | Val | Asn | His | Arg | Arg | Thr | His | Leu | Thr | Lys |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Leu | Met | His | Thr | Val | Glu | Gln | Ala | Thr | Leu | Arg | Ile | Ser | Gln | Ser | Phe |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Gln | Lys | Thr | Thr | Glu | Phe | Asp | Thr | Asn | Ser | Thr | Asp | Ile | Ala | Leu | Lys |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Val | Phe | Phe | Phe | Asp | Ser | Tyr | Asn | Met | Lys | His | Ile | His | Pro | His | Met |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Asn | Met | Asp | Gly | Asp | Tyr | Ile | Asn | Ile | Phe | Pro | Lys | Arg | Lys | Ala | Ala |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Tyr | Asp | Ser | Asn | Gly | Asn | Val | Ala | Val | Ala | Phe | Leu | Tyr | Tyr | Lys | Ser |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Ile | Gly | Pro | Leu | Leu | Ser | Ser | Ser | Asp | Asn | Phe | Leu | Leu | Lys | Pro | Gln |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Asn | Tyr | Asp | Asn | Ser | Glu | Glu | Glu | Glu | Arg | Val | Ile | Ser | Ser | Val | Ile |
| | | | 325 | | | | | | 330 | | | | | 335 | |
| Ser | Val | Ser | Met | Ser | Ser | Asn | Pro | Pro | Thr | Leu | Tyr | Glu | Leu | Glu | Lys |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| Ile | Thr | Phe | Thr | Leu | Ser | His | Arg | Lys | Val | Thr | Asp | Arg | Tyr | Arg | Ser |
| | | 355 | | | | | 360 | | | | | 365 | | | |
| Leu | Cys | Ala | Phe | Trp | Asn | Tyr | Ser | Pro | Asp | Thr | Met | Asn | Gly | Ser | Trp |
| | 370 | | | | | 375 | | | | | 380 | | | | |
| Ser | Ser | Glu | Gly | Cys | Glu | Leu | Thr | Tyr | Ser | Asn | Glu | Thr | His | Thr | Ser |
| 385 | | | | | 390 | | | | | 395 | | | | | 400 |
| Cys | Arg | Cys | Asn | His | Leu | Thr | His | Phe | Ala | Ile | Leu | Met | Ser | Ser | Gly |
| | | | 405 | | | | | 410 | | | | | | 415 | |
| Pro | Ser | Ile | Gly | Ile | Lys | Asp | Tyr | Asn | Ile | Leu | Thr | Arg | Ile | Thr | Gln |
| | | | 420 | | | | | 425 | | | | | 430 | | |
| Leu | Gly | Ile | Ile | Ile | Ser | Leu | Ile | Cys | Leu | Ala | Ile | Cys | Ile | Phe | Thr |
| | | 435 | | | | | 440 | | | | | 445 | | | |
| Phe | Trp | Phe | Phe | Ser | Glu | Ile | Gln | Ser | Thr | Arg | Thr | Thr | Ile | His | Lys |
| | 450 | | | | | 455 | | | | | 460 | | | | |
| Asn | Leu | Cys | Cys | Ser | Leu | Phe | Leu | Ala | Glu | Leu | Val | Phe | Leu | Val | Gly |
| 465 | | | | | 470 | | | | | 475 | | | | | 480 |
| Ile | Asn | Thr | Asn | Thr | Asn | Lys | Leu | Phe | Cys | Ser | Ile | Ile | Ala | Gly | Leu |

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<210> 51
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 51
ggtaatgagc tccattacag 20

<210> 52
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 52
ggagtagaaa gcgcatgg 18

<210> 53
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 53
cacctgatac catgaatggc ag 22

<210> 54
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 54
cgagctcgaa ttaattcg 18

<210> 55
<211> 18
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 55

ggatctcctg agctcagg

18

<210> 56

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 56

cctagttgag tgatccttgt aag

23

<210> 57

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 57

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50

<210> 58

<211> 2137

<212> DNA

<213> Homo sapiens

<400> 58

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gaaccctttc ccagcactt ggttttccaa catgatattt atgagtaatt tattttgata 2040
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<210> 59
 <211> 216
 <212> PRT
 <213> Homo sapiens

<400> 59

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| Met | Arg | Ser | Gly | Cys | Val | Val | Val | His | Val | Trp | Ile | Leu | Ala | Gly | Leu |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| | | | | | | | | | | | | | | | |
| Trp | Leu | Ala | Val | Ala | Gly | Arg | Pro | Leu | Ala | Phe | Ser | Asp | Ala | Gly | Pro |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| | | | | | | | | | | | | | | | |
| His | Val | His | Tyr | Gly | Trp | Gly | Asp | Pro | Ile | Arg | Leu | Arg | His | Leu | Tyr |
| | | | 35 | | | | 40 | | | | | 45 | | | |
| | | | | | | | | | | | | | | | |
| Thr | Ser | Gly | Pro | His | Gly | Leu | Ser | Ser | Cys | Phe | Leu | Arg | Ile | Arg | Ala |
| | | | 50 | | | 55 | | | | | 60 | | | | |
| | | | | | | | | | | | | | | | |
| Asp | Gly | Val | Val | Asp | Cys | Ala | Arg | Gly | Gln | Ser | Ala | His | Ser | Leu | Leu |
| 65 | | | | | 70 | | | | 75 | | | | | 80 | |
| | | | | | | | | | | | | | | | |
| Glu | Ile | Lys | Ala | Val | Ala | Leu | Arg | Thr | Val | Ala | Ile | Lys | Gly | Val | His |
| | | | | 85 | | | | 90 | | | | | | 95 | |
| | | | | | | | | | | | | | | | |
| Ser | Val | Arg | Tyr | Leu | Cys | Met | Gly | Ala | Asp | Gly | Lys | Met | Gln | Gly | Leu |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| | | | | | | | | | | | | | | | |
| Leu | Gln | Tyr | Ser | Glu | Glu | Asp | Cys | Ala | Phe | Glu | Glu | Glu | Ile | Arg | Pro |
| | | 115 | | | | 120 | | | | | | 125 | | | |
| | | | | | | | | | | | | | | | |
| Asp | Gly | Tyr | Asn | Val | Tyr | Arg | Ser | Glu | Lys | His | Arg | Leu | Pro | Val | Ser |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| | | | | | | | | | | | | | | | |
| Leu | Ser | Ser | Ala | Lys | Gln | Arg | Gln | Leu | Tyr | Lys | Asn | Arg | Gly | Phe | Leu |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| | | | | | | | | | | | | | | | |
| Pro | Leu | Ser | His | Phe | Leu | Pro | Met | Leu | Pro | Met | Val | Pro | Glu | Glu | Pro |
| | | | | 165 | | | | 170 | | | | | | 175 | |

Glu Asp Leu Arg Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu
180 185 190

Glu Thr Asp Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala
195 200 205

Val Arg Ser Pro Ser Phe Glu Lys
210 215

<210> 60
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 60
atccgcccag atggctacaa tgtgta 26

<210> 61
<211> 42
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 61
gcctcccggt ctccctgagc agtgccaaac agcggcagtg ta 42

<210> 62
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 62
ccagtccggt gacaagccca aa 22

<210> 63
<211> 1295
<212> DNA
<213> Homo sapiens

<400> 63
cccagaagtt caagggtccc cggcctcctg cgctcctgcc gccgggaccc tcgacctcct 60
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gctgctgctg cgctacctgg tggcgcctt gggctatcat aaggcctatg ggttttctgc 180
cccaaaagac caacaagtag tcacagcagt agagtaccaa gaggctattt tagcctgcaa 240
aaccctcaaag aagactgttt cctccagatt agagtggaag aaactgggtc ggagtgtctc 300

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ctttgtctac tatcaacaga ctcttcaagg tgattttaaa aatcgagctg agatgataga 360
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tagtgcccca tctgagcaag gccaaaacct ggaagaggat acagtcactc tggaagtatt 480
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tgtagaatc ttacaataaa tatagcttga tattc 1295

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<210> 64
 <211> 312
 <212> PRT
 <213> Homo sapiens

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<400> 64
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Leu Val Val Ala Leu Gly Tyr His Lys Ala Tyr Gly Phe Ser Ala Pro
      20              25              30

Lys Asp Gln Gln Val Val Thr Ala Val Glu Tyr Gln Glu Ala Ile Leu
      35              40              45

Ala Cys Lys Thr Pro Lys Lys Thr Val Ser Ser Arg Leu Glu Trp Lys
      50              55              60

Lys Leu Gly Arg Ser Val Ser Phe Val Tyr Tyr Gln Gln Thr Leu Gln
      65              70              75              80

Gly Asp Phe Lys Asn Arg Ala Glu Met Ile Asp Phe Asn Ile Arg Ile
      85              90              95

Lys Asn Val Thr Arg Ser Asp Ala Gly Lys Tyr Arg Cys Glu Val Ser
      100              105              110

Ala Pro Ser Glu Gln Gly Gln Asn Leu Glu Glu Asp Thr Val Thr Leu
      115              120              125

Glu Val Leu Val Ala Pro Ala Val Pro Ser Cys Glu Val Pro Ser Ser
      130              135              140

Ala Leu Ser Gly Thr Val Val Glu Leu Arg Cys Gln Asp Lys Glu Gly
      145              150              155              160

Asn Pro Ala Pro Glu Tyr Thr Trp Phe Lys Asp Gly Ile Arg Leu Leu
      165              170              175

```

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Asn | Pro | Arg | Leu | Gly | Ser | Gln | Ser | Thr | Asn | Ser | Ser | Tyr | Thr | Met |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Asn | Thr | Lys | Thr | Gly | Thr | Leu | Gln | Phe | Asn | Thr | Val | Ser | Lys | Leu | Asp |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Thr | Gly | Glu | Tyr | Ser | Cys | Glu | Ala | Arg | Asn | Ser | Val | Gly | Tyr | Arg | Arg |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Cys | Pro | Gly | Lys | Arg | Met | Gln | Val | Asp | Asp | Leu | Asn | Ile | Ser | Gly | Ile |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Ile | Ala | Ala | Val | Val | Val | Val | Ala | Leu | Val | Ile | Ser | Val | Cys | Gly | Leu |
| | | | 245 | | | | | | 250 | | | | | 255 | |
| Gly | Val | Cys | Tyr | Ala | Gln | Arg | Lys | Gly | Tyr | Phe | Ser | Lys | Glu | Thr | Ser |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Phe | Gln | Lys | Ser | Asn | Ser | Ser | Ser | Lys | Ala | Thr | Thr | Met | Ser | Glu | Asn |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Val | Gln | Trp | Leu | Thr | Pro | Val | Ile | Pro | Ala | Leu | Trp | Lys | Ala | Ala | Ala |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Gly | Gly | Ser | Arg | Gly | Gln | Glu | Phe | | | | | | | | |
| 305 | | | | | 310 | | | | | | | | | | |

<210> 65
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 65
 atcgttgtga agttagtgcc cc 22

<210> 66
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 66
 acctgcgata tccaacagaa ttg 23

<210> 67
 <211> 48
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 67

ggaagaggat acagtcactc tggaagtatt agtggctcca gcagttcc

48

<210> 68

<211> 2639

<212> DNA

<213> Homo sapiens

<400> 68

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| agaaagaaga | ggaagatggt | gggcaacatt | tatttaacat | gctccacagc | ccggaccctg | 120 |
| gcatcatgct | gctattcctg | caaatactga | agaagcatgg | gatttaaata | ttttacttct | 180 |
| aaataaatga | attactcaat | ctcctatgac | catctataca | tactccacct | tcaaaaagta | 240 |
| catcaatatt | atatcattaa | ggaaatagta | accttctctt | ctccaatatg | catgacattt | 300 |
| ttggacaatg | caattgtggc | actggcactt | atttcagtga | agaaaaactt | tgtggttcta | 360 |
| tggcattcat | catttgacaa | atgcaagcat | cttccttata | aatcagctcc | tattgaactt | 420 |
| actagcactg | actgtggaat | ccttaagggc | ccattacatt | tctgaagaag | aaagctaaga | 480 |
| tgaaggacat | gccactccga | attcatgtgc | tacttggcct | agctatcact | acactagtac | 540 |
| aagctgtaga | taaaaaagtg | gattgtccac | ggttatgtac | gtgtgaaatc | aggccttggg | 600 |
| ttacaccag | atccatttat | atggaagcat | ctacagtggg | ttgtaatgat | ttaggtcttt | 660 |
| taactttccc | agccagattg | ccagctaaca | cacagattct | tctcctacag | actaacaata | 720 |
| ttgcaaaaat | tgaatactcc | acagactttc | cagtaaacct | tactggcctg | gatttatctc | 780 |
| aaaacaattt | atcttcagtc | accaatatta | atgtaaaaaa | gatgcctcag | ctcctttctg | 840 |
| tgtacctaga | ggaaaacaaa | cttactgaac | tgcctgaaaa | atgtctgtcc | gaactgagca | 900 |
| acttacaaga | actctatatt | aatcacaaac | tgccttctac | aatttcacct | ggagccttta | 960 |
| ttggcctaca | taactttctt | cgacttcatc | tcaattcaaa | tagattgcag | atgatcaaca | 1020 |
| gtaagtgggt | tgatgtctct | ccaaatctag | agattctgat | gattggggaa | aatccaatta | 1080 |
| tcagaatcaa | agacatgaac | tttaagcctc | ttatcaatct | tcgcagcctg | gttatagctg | 1140 |
| gtataaacct | cacagaaata | ccagataacg | ccttggttgg | actggaaaac | ttagaaagca | 1200 |
| tctcttttta | cgataacagg | cttattaaag | taccccatgt | tgctcttcaa | aaagttgtaa | 1260 |
| atctcaaatt | tttgatctta | aataaaaatc | ctattaatag | aatacgaagg | ggtgatttta | 1320 |
| gcaatatgct | acacttaaaa | gagttgggga | taaataatat | gcctgagctg | atttccatcg | 1380 |
| atagtcttgc | tgtggataac | ctgccagatt | taagaaaaat | agaagctact | aacaacccta | 1440 |
| gattgtctta | cattcacccc | aatgcatttt | tcagactccc | caagctggaa | tcactcatgc | 1500 |
| tgaacagcaa | tgtctcagtc | gccctgtacc | atggtaccat | tgagtctctg | ccaaacctca | 1560 |
| aggaaatcag | catacacagt | aaccccatca | ggtgtgactg | tgtcatccgt | tggatgaaca | 1620 |
| tgaacaaaac | caacattcga | ttcatggagc | cagattcact | gttttgctg | gacccacctg | 1680 |
| aattccaagg | tcagaatggt | cggcaagtgc | atttcaggga | catgatggaa | atttgtctcc | 1740 |
| ctcttatagc | tcctgagagc | tttccttcta | atctaaatgt | agaagctggg | agctatgttt | 1800 |
| cctttcactg | tagagctact | gcagaaccac | agcctgaaat | ctactggata | acaccttctg | 1860 |
| gtcaaaaact | cttgccctaat | accctgacag | acaagttcta | tgtccattct | gagggaacac | 1920 |
| tagatataaa | tggcgtaact | cccaaagaag | ggggtttata | tacttgtata | gcaactaacc | 1980 |
| tagttggcgc | tgacttgaag | tctgttatga | tcaaagtgga | tggatctttt | ccacaagata | 2040 |
| acaatggctc | tttgaatatt | aaaataagag | atattcaggc | caattcagtt | ttggtgtcct | 2100 |
| ggaaagcaag | ttctaaaatt | ctcaaactta | gtgttaaatt | gacagccttt | gtcaagactg | 2160 |
| aaaattctca | tgtgcgcaa | agtgtctgaa | taccatctga | tgtcaaggta | tataatctta | 2220 |
| ctcatctgaa | tccatcaact | gagtataaaa | tttgtattga | tattcccacc | atctatcaga | 2280 |
| aaaacagaaa | aaaatgtgta | aatgtcacca | ccaaagggtt | gcaccctgat | caaaaagagt | 2340 |
| atgaaaagaa | taataccaca | acacttatgg | cctgtcttgg | aggccttctg | gggattattg | 2400 |
| gtgtgatatg | tcttatcagc | tgccctctct | cagaaatgaa | ctgtgatggg | ggacacagct | 2460 |
| atgtgaggaa | ttacttacag | aaaccaacct | ttgcattagg | tgagctttat | cctcctctga | 2520 |
| taaatctctg | ggaagcagga | aaagaaaaaa | gtacatcact | gaaagtaaaa | gcaactgtta | 2580 |
| taggtttacc | aacaaatatg | tcctaaaaac | caccaaggaa | acctactcca | aaaatgaac | 2639 |

<210> 69
 <211> 708
 <212> PRT
 <213> Homo sapiens

<400> 69
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 1 5 10 15
 Thr Thr Leu Val Gln Ala Val Asp Lys Lys Val Asp Cys Pro Arg Leu
 20 25 30
 Cys Thr Cys Glu Ile Arg Pro Trp Phe Thr Pro Arg Ser Ile Tyr Met
 35 40 45
 Glu Ala Ser Thr Val Asp Cys Asn Asp Leu Gly Leu Leu Thr Phe Pro
 50 55 60
 Ala Arg Leu Pro Ala Asn Thr Gln Ile Leu Leu Leu Gln Thr Asn Asn
 65 70 75 80
 Ile Ala Lys Ile Glu Tyr Ser Thr Asp Phe Pro Val Asn Leu Thr Gly
 85 90 95
 Leu Asp Leu Ser Gln Asn Asn Leu Ser Ser Val Thr Asn Ile Asn Val
 100 105 110
 Lys Lys Met Pro Gln Leu Leu Ser Val Tyr Leu Glu Glu Asn Lys Leu
 115 120 125
 Thr Glu Leu Pro Glu Lys Cys Leu Ser Glu Leu Ser Asn Leu Gln Glu
 130 135 140
 Leu Tyr Ile Asn His Asn Leu Leu Ser Thr Ile Ser Pro Gly Ala Phe
 145 150 155 160
 Ile Gly Leu His Asn Leu Leu Arg Leu His Leu Asn Ser Asn Arg Leu
 165 170 175
 Gln Met Ile Asn Ser Lys Trp Phe Asp Ala Leu Pro Asn Leu Glu Ile
 180 185 190
 Leu Met Ile Gly Glu Asn Pro Ile Ile Arg Ile Lys Asp Met Asn Phe
 195 200 205
 Lys Pro Leu Ile Asn Leu Arg Ser Leu Val Ile Ala Gly Ile Asn Leu
 210 215 220
 Thr Glu Ile Pro Asp Asn Ala Leu Val Gly Leu Glu Asn Leu Glu Ser
 225 230 235 240
 Ile Ser Phe Tyr Asp Asn Arg Leu Ile Lys Val Pro His Val Ala Leu
 245 250 255
 Gln Lys Val Val Asn Leu Lys Phe Leu Asp Leu Asn Lys Asn Pro Ile
 260 265 270

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Arg | Ile | Arg | Arg | Gly | Asp | Phe | Ser | Asn | Met | Leu | His | Leu | Lys | Glu |
| | | 275 | | | | 280 | | | | | | 285 | | | |
| Leu | Gly | Ile | Asn | Asn | Met | Pro | Glu | Leu | Ile | Ser | Ile | Asp | Ser | Leu | Ala |
| | | 290 | | | | 295 | | | | | 300 | | | | |
| Val | Asp | Asn | Leu | Pro | Asp | Leu | Arg | Lys | Ile | Glu | Ala | Thr | Asn | Asn | Pro |
| | | 305 | | | 310 | | | | | 315 | | | | | |
| Arg | Leu | Ser | Tyr | Ile | His | Pro | Asn | Ala | Phe | Phe | Arg | Leu | Pro | Lys | Leu |
| | | | | 325 | | | | | 330 | | | | | | |
| Glu | Ser | Leu | Met | Leu | Asn | Ser | Asn | Ala | Leu | Ser | Ala | Leu | Tyr | His | Gly |
| | | | 340 | | | | | 345 | | | | | | | |
| Thr | Ile | Glu | Ser | Leu | Pro | Asn | Leu | Lys | Glu | Ile | Ser | Ile | His | Ser | Asn |
| | | 355 | | | | 360 | | | | | | 365 | | | |
| Pro | Ile | Arg | Cys | Asp | Cys | Val | Ile | Arg | Trp | Met | Asn | Met | Asn | Lys | Thr |
| | | 370 | | | | 375 | | | | | 380 | | | | |
| Asn | Ile | Arg | Phe | Met | Glu | Pro | Asp | Ser | Leu | Phe | Cys | Val | Asp | Pro | Pro |
| | | 385 | | | 390 | | | | | 395 | | | | | |
| Glu | Phe | Gln | Gly | Gln | Asn | Val | Arg | Gln | Val | His | Phe | Arg | Asp | Met | Met |
| | | | | 405 | | | | | 410 | | | | | | |
| Glu | Ile | Cys | Leu | Pro | Leu | Ile | Ala | Pro | Glu | Ser | Phe | Pro | Ser | Asn | Leu |
| | | | 420 | | | | | 425 | | | | | | | |
| Asn | Val | Glu | Ala | Gly | Ser | Tyr | Val | Ser | Phe | His | Cys | Arg | Ala | Thr | Ala |
| | | 435 | | | | | 440 | | | | 445 | | | | |
| Glu | Pro | Gln | Pro | Glu | Ile | Tyr | Trp | Ile | Thr | Pro | Ser | Gly | Gln | Lys | Leu |
| | | 450 | | | | 455 | | | | | 460 | | | | |
| Leu | Pro | Asn | Thr | Leu | Thr | Asp | Lys | Phe | Tyr | Val | His | Ser | Glu | Gly | Thr |
| | | 465 | | | 470 | | | | | 475 | | | | | |
| Leu | Asp | Ile | Asn | Gly | Val | Thr | Pro | Lys | Glu | Gly | Gly | Leu | Tyr | Thr | Cys |
| | | | | 485 | | | | | 490 | | | | | | |
| Ile | Ala | Thr | Asn | Leu | Val | Gly | Ala | Asp | Leu | Lys | Ser | Val | Met | Ile | Lys |
| | | | 500 | | | | | 505 | | | | 510 | | | |
| Val | Asp | Gly | Ser | Phe | Pro | Gln | Asp | Asn | Asn | Gly | Ser | Leu | Asn | Ile | Lys |
| | | 515 | | | | | 520 | | | | | 525 | | | |
| Ile | Arg | Asp | Ile | Gln | Ala | Asn | Ser | Val | Leu | Val | Ser | Trp | Lys | Ala | Ser |
| | | 530 | | | | 535 | | | | | 540 | | | | |
| Ser | Lys | Ile | Leu | Lys | Ser | Ser | Val | Lys | Trp | Thr | Ala | Phe | Val | Lys | Thr |
| | | 545 | | | 550 | | | | | 555 | | | | | |
| Glu | Asn | Ser | His | Ala | Ala | Gln | Ser | Ala | Arg | Ile | Pro | Ser | Asp | Val | Lys |
| | | | | 565 | | | | | 570 | | | | | | |

Val Tyr Asn Leu Thr His Leu Asn Pro Ser Thr Glu Tyr Lys Ile Cys
580 585 590

Ile Asp Ile Pro Thr Ile Tyr Gln Lys Asn Arg Lys Lys Cys Val Asn
595 600 605

Val Thr Thr Lys Gly Leu His Pro Asp Gln Lys Glu Tyr Glu Lys Asn
610 615 620

Asn Thr Thr Thr Leu Met Ala Cys Leu Gly Gly Leu Leu Gly Ile Ile
625 630 635 640

Gly Val Ile Cys Leu Ile Ser Cys Leu Ser Pro Glu Met Asn Cys Asp
645 650 655

Gly Gly His Ser Tyr Val Arg Asn Tyr Leu Gln Lys Pro Thr Phe Ala
660 665 670

Leu Gly Glu Leu Tyr Pro Pro Leu Ile Asn Leu Trp Glu Ala Gly Lys
675 680 685

Glu Lys Ser Thr Ser Leu Lys Val Lys Ala Thr Val Ile Gly Leu Pro
690 695 700

Thr Asn Met Ser
705

<210> 70
<211> 1305
<212> DNA
<213> Homo sapiens

<400> 70
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agctgcagcc ttttgaaaca cgcaagaagg aaatcaatag tgtggacagg gctggaacct 120
ttaccacgct tgttgagta gatgaggaat gggctcgtga ttatgctgac attccagcat 180
gaatctggta gacctgtggt taaccggttc cctctccatg tgtctcctcc tacaaggtt 240
tgttcttatg atactgtgct ttcattctgc cagtatgtgt cccaagggtt gtctttgttc 300
ttcctctggg ggtttaaatg tcacctgtag caatgcaaat ctcaaggaaa tacctagaga 360
tcttctcct gaaacagtct tactgtatct ggactccaat cagatcacat ctattcccaa 420
tgaaatttt aaggacctcc atcaactgag agttctcaac ctgtccaaaa atggcattga 480
gtttatcgat gagcatgcct tcaaaggagt agctgaaacc ttgcagactc tggacttgtc 540
cgacaatcgg attcaaagtg tgcacaaaaa tgccttcaat aacctgaagg ccagggccag 600
aattgccaac aacccttggc actgcgactg tactctacag caagttctga ggagcatggc 660
gtccaatcat gagacagccc acaacgtgat ctgtaaaacg tccgtgttgg atgaacatgc 720
tggcagacca ttctcaatg ctgccaacga cgctgacctt tgtaacctcc ctaaaaaaac 780
taccgattat gccatgctgg tcaccatggt tggctgggtc actatggtga tctcatatgt 840
ggatatattat gtgaggcaaa atcaggagga tgcccggaga cacctcgaat acttgaaatc 900
cctgccaaagc aggcagaaga aagcagatga acctgatgat attagcactg tggatatagt 960
tccaaactga ctgtcattga gaaagaaaga aagtagtttg cgattgcagt agaaataagt 1020
ggtttacttc tcccatccat tgtaaacatt tgaaactttg tatttcagtt tttttgaaat 1080
tatgccactg ctgaactttt aacaaacact acaacataaa taatttgagt ttaggtgatc 1140
cacccttaa ttgtaccccc gatggtatat ttctgagtaa gctactatct gaacattagt 1200
tagatccatc tcaactattt ataatgaaat ttattttttt aattttaaaag caaataaaaag 1260
cttaactttg aaccatggga aaaaaaaaaa aaaaaaaaaa aaaca 1305

<210> 71

<211> 259
<212> PRT
<213> Homo sapiens

<400> 71

| | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Met | Asn | Leu | Val | Asp | Leu | Trp | Leu | Thr | Arg | Ser | Leu | Ser | Met | Cys | Leu | |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | | |
| Leu | Leu | Gln | Ser | Phe | Val | Leu | Met | Ile | Leu | Cys | Phe | His | Ser | Ala | Ser | |
| | | | 20 | | | | | 25 | | | | | 30 | | | |
| Met | Cys | Pro | Lys | Gly | Cys | Leu | Cys | Ser | Ser | Ser | Gly | Gly | Leu | Asn | Val | |
| | | 35 | | | | | 40 | | | | | 45 | | | | |
| Thr | Cys | Ser | Asn | Ala | Asn | Leu | Lys | Glu | Ile | Pro | Arg | Asp | Leu | Pro | Pro | |
| | 50 | | | | | 55 | | | | | 60 | | | | | |
| Glu | Thr | Val | Leu | Leu | Tyr | Leu | Asp | Ser | Asn | Gln | Ile | Thr | Ser | Ile | Pro | |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 | |
| Asn | Glu | Ile | Phe | Lys | Asp | Leu | His | Gln | Leu | Arg | Val | Leu | Asn | Leu | Ser | |
| | | | | 85 | | | | | 90 | | | | | | 95 | |
| Lys | Asn | Gly | Ile | Glu | Phe | Ile | Asp | Glu | His | Ala | Phe | Lys | Gly | Val | Ala | |
| | | | 100 | | | | | 105 | | | | | 110 | | | |
| Glu | Thr | Leu | Gln | Thr | Leu | Asp | Leu | Ser | Asp | Asn | Arg | Ile | Gln | Ser | Val | |
| | | 115 | | | | | 120 | | | | | | 125 | | | |
| His | Lys | Asn | Ala | Phe | Asn | Asn | Leu | Lys | Ala | Arg | Ala | Arg | Ile | Ala | Asn | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Asn | Pro | Trp | His | Cys | Asp | Cys | Thr | Leu | Gln | Gln | Val | Leu | Arg | Ser | Met | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | |
| Ala | Ser | Asn | His | Glu | Thr | Ala | His | Asn | Val | Ile | Cys | Lys | Thr | Ser | Val | |
| | | | | 165 | | | | | 170 | | | | | 175 | | |
| Leu | Asp | Glu | His | Ala | Gly | Arg | Pro | Phe | Leu | Asn | Ala | Ala | Asn | Asp | Ala | |
| | | | 180 | | | | | 185 | | | | | | 190 | | |
| Asp | Leu | Cys | Asn | Leu | Pro | Lys | Lys | Thr | Thr | Asp | Tyr | Ala | Met | Leu | Val | |
| | | 195 | | | | | 200 | | | | | 205 | | | | |
| Thr | Met | Phe | Gly | Trp | Phe | Thr | Met | Val | Ile | Ser | Tyr | Val | Val | Tyr | Tyr | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Val | Arg | Gln | Asn | Gln | Glu | Asp | Ala | Arg | Arg | His | Leu | Glu | Tyr | Leu | Lys | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | |
| Ser | Leu | Pro | Ser | Arg | Gln | Lys | Lys | Ala | Asp | Glu | Pro | Asp | Asp | Ile | Ser | |
| | | | | 245 | | | | | 250 | | | | | | 255 | |
| Thr | Val | Val | | | | | | | | | | | | | | |

<210> 72

<211> 2290
 <212> DNA
 <213> Homo sapiens

<400> 72
 accgagccga gcggaccgaa ggcgcgcccc agatgcaggt gagcaagagg atgctggcgg 60
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 tgctgggctc agtgctgtca ggctcggcca cgggctgccc gccccgctgc gagggtccg 180
 cccaggaccg cgctgtgctg tgccaccgca agtgctttgt ggaggtcccc gagggcatcc 240
 ccaccgagac gcgcctgctg gacctaggca agaaccgcat caaacgctc aaccaggacg 300
 agttcgccag cttcccgac ctggaggagc tggagctcaa cgagaacatc gtgagcgccg 360
 tggagccccg cgccttcaac aacctcttca acctccggac gctgggtctc cgcagcaacc 420
 gcctgaagct catcccgcta ggctgttca ctggcctcag caacctgacc aagcaggaca 480
 tcagcgagaa caagatcggt atcctactgg actacatggt tcaggacctg tacaacctca 540
 agtcaactgga ggttggcgac aatgacctcg tctacatctc tcaccgcgcc ttcagcggcc 600
 tcaacagcct ggagcagctg acgctggaga aatgcaacct gacctccatc cccaccgagg 660
 cgctgtccca cctgcacggc ctcatcgctc tgaggtccg gcacctcaac atcaatgcca 720
 tccgggacta ctcttcaag aggtgttacc gactcaaggt cttggagatc tccactggc 780
 cctacttga caccatgaca cccaactgcc tctacggcct caacctgacg tccctgtcca 840
 tcacacactg caatctgacc gctgtgccct acctggcgt ccgccacctg gtctatctcc 900
 gcttctctcaa cctctctac aaccccatca gcaccattga gggctccatg ttgcatgagc 960
 tgctccggct gcaggagatc cagctggtgg gcgggcagct ggccgtggtg gagccctatg 1020
 ccttccgcgg cctcaactac ctgcgcgtgc tcaatgtctc tggcaaccag ctgaccacac 1080
 tggaggaatc agtcttccac tcgggtgggca acctggagac actcatcttg gactccaacc 1140
 cgctggcctg cgactgtcgg ctctgtggg tggtccggcg ccgctggcgg ctcaacttca 1200
 accggcagca gccacgtgc gccacgccc agtttgtcca gggcaaggag ttcaaggact 1260
 tccctgatgt gctactgccc aactacttca cctgccgcgg cgcccgcatc cgggaccgca 1320
 aggcccagca ggtgtttgtg gacgagggcc acacggtgca gtttgtgtgc cgggcccgatg 1380
 gcgacccgcc gcccgccatc ctctggctct caccgccaaa gcacctggtc tcagccaaga 1440
 gcaatgggcg gctcacagtc ttccctgatg gcacgtgga ggtgcgtac gccagggtac 1500
 aggacaacgg cagctacctg tgcatcgcg ccaacgcggg cggcaacgac tccatgcccg 1560
 cccacctgca tgtgcgcagc tactcgccc actggcccca tcagcccaac aagaccttcg 1620
 cttcatctc caaccagccg ggcgaggag aggccaaacag caccgcgcc actgtgcctt 1680
 tcccttgcga catcaagacc ctcatcatcg ccaccaccat gggcttcatc tcttctctgg 1740
 gcgtcgctct cttctgcctg gtgctgctgt ttctctggag ccggggcaag ggcaacacaa 1800
 agcacaacat cgagatcgag tatgtgcccc gaaagtgcga cgcaggcatc agctccgcgg 1860
 acgcgccccg caagttcaac atgaagatga tatgaggccg gggcgggggg cagggacccc 1920
 cgggcggccg ggcaggggaa ggggcctggt cgccacctgc tcaactctca gtccttccca 1980
 cctcctccct acccttctac acacgttctc tttctccctc ccgcctccgt cccctgctgc 2040
 cccccgccag ccctcaccac ctgccctcct tctaccagga cctcagaagc ccagacctgg 2100
 ggacccacc tacacagggg cattgacaga ctggagtga aagccgacga accgacacgc 2160
 ggcagagtca ataattcaat aaaaaagtta cgaactttct ctgtaacttg ggtttcaata 2220
 attatggatt tttatgaaaa cttgaaataa taaaaagaga aaaaaactaa aaaaaaaaaa 2280
 aaaaaaaaaa 2290

<210> 73
 <211> 620
 <212> PRT
 <213> Homo sapiens

<400> 73
 Met Gln Val Ser Lys Arg Met Leu Ala Gly Gly Val Arg Ser Met Pro
 1 5 10 15
 Ser Pro Leu Leu Ala Cys Trp Gln Pro Ile Leu Leu Leu Val Leu Gly
 20 25 30

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Val | Leu | Ser | Gly | Ser | Ala | Thr | Gly | Cys | Pro | Pro | Arg | Cys | Glu | Cys | 35 | 40 | 45 |
| Ser | Ala | Gln | Asp | Arg | Ala | Val | Leu | Cys | His | Arg | Lys | Cys | Phe | Val | Ala | 50 | 55 | 60 |
| Val | Pro | Glu | Gly | Ile | Pro | Thr | Glu | Thr | Arg | Leu | Leu | Asp | Leu | Gly | Lys | 65 | 70 | 75 |
| Asn | Arg | Ile | Lys | Thr | Leu | Asn | Gln | Asp | Glu | Phe | Ala | Ser | Phe | Pro | His | 85 | 90 | 95 |
| Leu | Glu | Glu | Leu | Glu | Leu | Asn | Glu | Asn | Ile | Val | Ser | Ala | Val | Glu | Pro | 100 | 105 | 110 |
| Gly | Ala | Phe | Asn | Asn | Leu | Phe | Asn | Leu | Arg | Thr | Leu | Gly | Leu | Arg | Ser | 115 | 120 | 125 |
| Asn | Arg | Leu | Lys | Leu | Ile | Pro | Leu | Gly | Val | Phe | Thr | Gly | Leu | Ser | Asn | 130 | 135 | 140 |
| Leu | Thr | Lys | Gln | Asp | Ile | Ser | Glu | Asn | Lys | Ile | Val | Ile | Leu | Leu | Asp | 145 | 150 | 155 |
| Tyr | Met | Phe | Gln | Asp | Leu | Tyr | Asn | Leu | Lys | Ser | Leu | Glu | Val | Gly | Asp | 165 | 170 | 175 |
| Asn | Asp | Leu | Val | Tyr | Ile | Ser | His | Arg | Ala | Phe | Ser | Gly | Leu | Asn | Ser | 180 | 185 | 190 |
| Leu | Glu | Gln | Leu | Thr | Leu | Glu | Lys | Cys | Asn | Leu | Thr | Ser | Ile | Pro | Thr | 195 | 200 | 205 |
| Glu | Ala | Leu | Ser | His | Leu | His | Gly | Leu | Ile | Val | Leu | Arg | Leu | Arg | His | 210 | 215 | 220 |
| Leu | Asn | Ile | Asn | Ala | Ile | Arg | Asp | Tyr | Ser | Phe | Lys | Arg | Leu | Tyr | Arg | 225 | 230 | 235 |
| Leu | Lys | Val | Leu | Glu | Ile | Ser | His | Trp | Pro | Tyr | Leu | Asp | Thr | Met | Thr | 245 | 250 | 255 |
| Pro | Asn | Cys | Leu | Tyr | Gly | Leu | Asn | Leu | Thr | Ser | Leu | Ser | Ile | Thr | His | 260 | 265 | 270 |
| Cys | Asn | Leu | Thr | Ala | Val | Pro | Tyr | Leu | Ala | Val | Arg | His | Leu | Val | Tyr | 275 | 280 | 285 |
| Leu | Arg | Phe | Leu | Asn | Leu | Ser | Tyr | Asn | Pro | Ile | Ser | Thr | Ile | Glu | Gly | 290 | 295 | 300 |
| Ser | Met | Leu | His | Glu | Leu | Leu | Arg | Leu | Gln | Glu | Ile | Gln | Leu | Val | Gly | 305 | 310 | 315 |
| Gly | Gln | Leu | Ala | Val | Val | Glu | Pro | Tyr | Ala | Phe | Arg | Gly | Leu | Asn | Tyr | 325 | 330 | 335 |

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 74

tcacctggag cctttattgg cc

22

<210> 75

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 75

ataccagcta taaccaggct gcg

23

<210> 76

<211> 52

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 76

caacagtaag tggtttgatg ctcttccaaa tctagagatt ctgatgattg
gg

50

52

<210> 77

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 77

ccatgtgtct cctcctacaa ag

22

<210> 78

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 78

gggaatagat gtgatctgat tgg

23

<210> 79
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 79
cacctgtagc aatgcaaatc tcaaggaaat acctagagat cttcctcctg 50

<210> 80
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 80
agcaaccgcc tgaagctcat cc 22

<210> 81
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 81
aaggcgcggt gaaagatgta gacg 24

<210> 82
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 82
gactacatgt ttcaggacct gtacaacctc aagtcactgg aggttggcga 50

<210> 83
<211> 1685
<212> DNA
<213> Homo sapiens

<400> 83
cccacgcgtc cgcacctcgg ccccgggctc cgaagcggct cgggggccc ctttcggtca 60
acatcgtagt ccacccctc cccatcccca gccccgggg attcaggctc gccagcgccc 120

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agccagggag cgggccggga agcgcgatgg gggccccagc cgcctcgctc ctgctcctgc 180
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<210> 84
 <211> 398
 <212> PRT
 <213> Homo sapiens

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<400> 84
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Pro Trp Thr Ser Asp Glu Thr Val Val Ala Gly Gly Thr Val Val Leu
      35                      40                      45

Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp Ser Asn
      50                      55                      60

Pro Ala Gln Gln Thr Leu Tyr Phe Gly Glu Lys Arg Ala Leu Arg Asp
      65                      70                      75                      80

Asn Arg Ile Gln Leu Val Thr Ser Thr Pro His Glu Leu Ser Ile Ser
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Ile Ser Asn Val Ala Leu Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile
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Phe Thr Met Pro Val Arg Thr Ala Lys Ser Leu Val Thr Val Leu Gly
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| Lys | Asp | Thr | Ala | Thr | Leu | Asn | Cys | Gln | Ser | Ser | Gly | Ser | Lys | Pro | Ala | | |
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| Ser | Val | Thr | Phe | Gln | Val | Thr | Arg | Glu | Asp | Asp | Gly | Ala | Ser | Ile | Val | | |
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| Cys | Ser | Val | Asn | His | Glu | Ser | Leu | Lys | Gly | Ala | Asp | Arg | Ser | Thr | Ser | | |
| | 210 | | | | | 215 | | | | | 220 | | | | | | |
| Gln | Arg | Ile | Glu | Val | Leu | Tyr | Thr | Pro | Thr | Ala | Met | Ile | Arg | Pro | Asp | | |
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| Pro | Pro | His | Pro | Arg | Glu | Gly | Gln | Lys | Leu | Leu | Leu | His | Cys | Glu | Gly | | |
| | | | | 245 | | | | | 250 | | | | | 255 | | | |
| Arg | Gly | Asn | Pro | Val | Pro | Gln | Gln | Tyr | Leu | Trp | Glu | Lys | Glu | Gly | Ser | | |
| | | | 260 | | | | | 265 | | | | | 270 | | | | |
| Val | Pro | Pro | Leu | Lys | Met | Thr | Gln | Glu | Ser | Ala | Leu | Ile | Phe | Pro | Phe | | |
| | | 275 | | | | | 280 | | | | | 285 | | | | | |
| Leu | Asn | Lys | Ser | Asp | Ser | Gly | Thr | Tyr | Gly | Cys | Thr | Ala | Thr | Ser | Asn | | |
| | 290 | | | | | 295 | | | | | 300 | | | | | | |
| Met | Gly | Ser | Tyr | Lys | Ala | Tyr | Tyr | Thr | Leu | Asn | Val | Asn | Asp | Pro | Ser | | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | | |
| Pro | Val | Pro | Ser | Ser | Ser | Ser | Thr | Tyr | His | Ala | Ile | Ile | Gly | Gly | Ile | | |
| | | | | 325 | | | | | 330 | | | | | 335 | | | |
| Val | Ala | Phe | Ile | Val | Phe | Leu | Leu | Leu | Ile | Met | Leu | Ile | Phe | Leu | Gly | | |
| | | | 340 | | | | | 345 | | | | | 350 | | | | |
| His | Tyr | Leu | Ile | Arg | His | Lys | Gly | Thr | Tyr | Leu | Thr | His | Glu | Ala | Lys | | |
| | | 355 | | | | | 360 | | | | | 365 | | | | | |
| Gly | Ser | Asp | Asp | Ala | Pro | Asp | Ala | Asp | Thr | Ala | Ile | Ile | Asn | Ala | Glu | | |
| | 370 | | | | | 375 | | | | | 380 | | | | | | |
| Gly | Gly | Gln | Ser | Gly | Gly | Asp | Asp | Lys | Lys | Glu | Tyr | Phe | Ile | | | | |
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<210> 85

<211> 22

<212> DNA

<213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 85
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<210> 86
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 86
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<210> 87
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 87
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<210> 88
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 88
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<210> 89
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

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<210> 90
 <211> 2755

<212> DNA
<213> Homo sapiens

<400> 90

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<210> 91
<211> 696
<212> PRT
<213> Homo sapiens

<400> 91

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| 1 | | | | 5 | | | | | 10 | | | | | 15 | | |
| Gly | Asn | Val | Thr | Gly | Asp | Val | Cys | Lys | Glu | Lys | Ile | Cys | Ser | Cys | Asn | |
| | | | 20 | | | | | 25 | | | | | 30 | | | |
| Glu | Ile | Glu | Gly | Asp | Leu | His | Val | Asp | Cys | Glu | Lys | Lys | Gly | Phe | Thr | |
| | | 35 | | | | | 40 | | | | | 45 | | | | |
| Ser | Leu | Gln | Arg | Phe | Thr | Ala | Pro | Thr | Ser | Gln | Phe | Tyr | His | Leu | Phe | |
| | 50 | | | | | 55 | | | | | 60 | | | | | |
| Leu | His | Gly | Asn | Ser | Leu | Thr | Arg | Leu | Phe | Pro | Asn | Glu | Phe | Ala | Asn | |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 | |
| Phe | Tyr | Asn | Ala | Val | Ser | Leu | His | Met | Glu | Asn | Asn | Gly | Leu | His | Glu | |
| | | | | 85 | | | | | 90 | | | | | 95 | | |
| Ile | Val | Pro | Gly | Ala | Phe | Leu | Gly | Leu | Gln | Leu | Val | Lys | Arg | Leu | His | |
| | | 100 | | | | | | 105 | | | | | 110 | | | |
| Ile | Asn | Asn | Asn | Lys | Ile | Lys | Ser | Phe | Arg | Lys | Gln | Thr | Phe | Leu | Gly | |
| | 115 | | | | | | 120 | | | | | 125 | | | | |
| Leu | Asp | Asp | Leu | Glu | Tyr | Leu | Gln | Ala | Asp | Phe | Asn | Leu | Leu | Arg | Asp | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Ile | Asp | Pro | Gly | Ala | Phe | Gln | Asp | Leu | Asn | Lys | Leu | Glu | Val | Leu | Ile | |
| 145 | | | | 150 | | | | | 155 | | | | | | 160 | |
| Leu | Asn | Asp | Asn | Leu | Ile | Ser | Thr | Leu | Pro | Ala | Asn | Val | Phe | Gln | Tyr | |
| | | | 165 | | | | | 170 | | | | | | 175 | | |
| Val | Pro | Ile | Thr | His | Leu | Asp | Leu | Arg | Gly | Asn | Arg | Leu | Lys | Thr | Leu | |
| | | 180 | | | | | | 185 | | | | | 190 | | | |
| Pro | Tyr | Glu | Glu | Val | Leu | Glu | Gln | Ile | Pro | Gly | Ile | Ala | Glu | Ile | Leu | |
| | 195 | | | | | | 200 | | | | | 205 | | | | |
| Leu | Glu | Asp | Asn | Pro | Trp | Asp | Cys | Thr | Cys | Asp | Leu | Leu | Ser | Leu | Lys | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Glu | Trp | Leu | Glu | Asn | Ile | Pro | Lys | Asn | Ala | Leu | Ile | Gly | Arg | Val | Val | |
| 225 | | | | 230 | | | | | | 235 | | | | | 240 | |
| Cys | Glu | Ala | Pro | Thr | Arg | Leu | Gln | Gly | Lys | Asp | Leu | Asn | Glu | Thr | Thr | |
| | | | 245 | | | | | | 250 | | | | | 255 | | |
| Glu | Gln | Asp | Leu | Cys | Pro | Leu | Lys | Asn | Arg | Val | Asp | Ser | Ser | Leu | Pro | |
| | | 260 | | | | | | 265 | | | | | 270 | | | |
| Ala | Pro | Pro | Ala | Gln | Glu | Glu | Thr | Phe | Ala | Pro | Gly | Pro | Leu | Pro | Thr | |
| | 275 | | | | | | 280 | | | | | 285 | | | | |
| Pro | Phe | Lys | Thr | Asn | Gly | Gln | Glu | Asp | His | Ala | Thr | Pro | Gly | Ser | Ala | |
| | 290 | | | | | 295 | | | | | 300 | | | | | |

Pro Asn Gly Gly Thr Lys Ile Pro Gly Asn Trp Gln Ile Lys Ile Arg
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 Pro Thr Ala Ala Ile Ala Thr Gly Ser Ser Arg Asn Lys Pro Leu Ala
 325 330 335
 Asn Ser Leu Pro Cys Pro Gly Gly Cys Ser Cys Asp His Ile Pro Gly
 340 345 350
 Ser Gly Leu Lys Met Asn Cys Asn Asn Arg Asn Val Ser Ser Leu Ala
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 Asp Leu Lys Pro Lys Leu Ser Asn Val Gln Glu Leu Phe Leu Arg Asp
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 Asn Lys Ile His Ser Ile Arg Lys Ser His Phe Val Asp Tyr Lys Asn
 385 390 395 400
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 Gly Val Leu Asp Gln Leu Thr Ser Ile Ile Gln Ile Asp Leu His Gly
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 Ser Lys Asn Ser Thr Gly Leu Ala Glu Thr Gly Thr His Ser Asn Ser
 595 600 605

Tyr Leu Asp Thr Ser Arg Val Ser Ile Ser Val Leu Val Pro Gly Leu
610 615 620

Leu Leu Val Phe Val Thr Ser Ala Phe Thr Val Val Gly Met Leu Val
625 630 635 640

Phe Ile Leu Arg Asn Arg Lys Arg Ser Lys Arg Arg Asp Ala Asn Ser
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Ser Ala Ser Glu Ile Asn Ser Leu Gln Thr Val Cys Asp Ser Ser Tyr
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Trp His Asn Gly Pro Tyr Asn Ala Asp Gly Ala His Arg Val Tyr Asp
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Cys Gly Ser His Ser Leu Ser Asp
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<210> 92
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 92
gttgatctg ggcaacaata ac 22

<210> 93
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 93
attgttgatgc aggctgagtt taag 24

<210> 94
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 94
ggtggctata catggatagc aattacctgg acacgctgtc ccggg 45

<210> 95
<211> 2226
<212> DNA

<213> Homo sapiens

<400> 95

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<210> 96

<211> 490

<212> PRT

<213> Homo sapiens

<400> 96

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Ala Ser Gly Ala Cys Tyr Ser Leu His His Ala Thr Met Lys Arg Gln
  35                      40                     45
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Ala Ala Glu Glu Ala Cys Ile Leu Arg Gly Gly Ala Leu Ser Thr Val
 50 55 60

Arg Ala Gly Ala Glu Leu Arg Ala Val Leu Ala Leu Leu Arg Ala Gly
 65 70 75 80

Pro Gly Pro Gly Gly Gly Ser Lys Asp Leu Leu Phe Trp Val Ala Leu
 85 90 95

Glu Arg Arg Arg Ser His Cys Thr Leu Glu Asn Glu Pro Leu Arg Gly
 100 105 110

Phe Ser Trp Leu Ser Ser Asp Pro Gly Gly Leu Glu Ser Asp Thr Leu
 115 120 125

Gln Trp Val Glu Glu Pro Gln Arg Ser Cys Thr Ala Arg Arg Cys Ala
 130 135 140

Val Leu Gln Ala Thr Gly Gly Val Glu Pro Ala Gly Trp Lys Glu Met
 145 150 155 160

Arg Cys His Leu Arg Ala Asn Gly Tyr Leu Cys Lys Tyr Gln Phe Glu
 165 170 175

Val Leu Cys Pro Ala Pro Arg Pro Gly Ala Ala Ser Asn Leu Ser Tyr
 180 185 190

Arg Ala Pro Phe Gln Leu His Ser Ala Ala Leu Asp Phe Ser Pro Pro
 195 200 205

Gly Thr Glu Val Ser Ala Leu Cys Arg Gly Gln Leu Pro Ile Ser Val
 210 215 220

Thr Cys Ile Ala Asp Glu Ile Gly Ala Arg Trp Asp Lys Leu Ser Gly
 225 230 235 240

Asp Val Leu Cys Pro Cys Pro Gly Arg Tyr Leu Arg Ala Gly Lys Cys
 245 250 255

Ala Glu Leu Pro Asn Cys Leu Asp Asp Leu Gly Gly Phe Ala Cys Glu
 260 265 270

Cys Ala Thr Gly Phe Glu Leu Gly Lys Asp Gly Arg Ser Cys Val Thr
 275 280 285

Ser Gly Glu Gly Gln Pro Thr Leu Gly Gly Thr Gly Val Pro Thr Arg
 290 295 300

Arg Pro Pro Ala Thr Ala Thr Ser Pro Val Pro Gln Arg Thr Trp Pro
 305 310 315 320

Ile Arg Val Asp Glu Lys Leu Gly Glu Thr Pro Leu Val Pro Glu Gln
 325 330 335

Asp Asn Ser Val Thr Ser Ile Pro Glu Ile Pro Arg Trp Gly Ser Gln
 340 345 350

Ser Thr Met Ser Thr Leu Gln Met Ser Leu Gln Ala Glu Ser Lys Ala
 355 360 365
 Thr Ile Thr Pro Ser Gly Ser Val Ile Ser Lys Phe Asn Ser Thr Thr
 370 375 380
 Ser Ser Ala Thr Pro Gln Ala Phe Asp Ser Ser Ser Ala Val Val Phe
 385 390 395 400
 Ile Phe Val Ser Thr Ala Val Val Val Leu Val Ile Leu Thr Met Thr
 405 410 415
 Val Leu Gly Leu Val Lys Leu Cys Phe His Glu Ser Pro Ser Ser Gln
 420 425 430
 Pro Arg Lys Glu Ser Met Gly Pro Pro Gly Leu Glu Ser Asp Pro Glu
 435 440 445
 Pro Ala Ala Leu Gly Ser Ser Ser Ala His Cys Thr Asn Asn Gly Val
 450 455 460
 Lys Val Gly Asp Cys Asp Leu Arg Asp Arg Ala Glu Gly Ala Leu Leu
 465 470 475 480
 Ala Glu Ser Pro Leu Gly Ser Ser Asp Ala
 485 490

<210> 97

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 97

tggaaggaga tgcgatgccca cctg

24

<210> 98

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 98

tgaccagtgg ggaaggacag

20

<210> 99

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 99
acagagcaga ggggtgccttg 20

<210> 100
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 100
tcagggacaa gtggtgtctc tccc 24

<210> 101
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 101
tcagggaagg agtgtgcagt tctg 24

<210> 102
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 102
acagctccccg atctcagtta cttgcatcgc ggacgaaatc ggcgctcgct 50

<210> 103
<211> 2026
<212> DNA
<213> Homo sapiens

<400> 103
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tatcccccg ctacctgggc cgccccgcgg cgggtgcgcg gtgagaggga gcgcgcgggc 180
agccgagcgc cgggtgtgagc cagcgtgtgt gccagtgtga gcggcggtgt gagcgcggtg 240
ggtgcggagg ggcgtgtgtg ccggcgcgcg cgccgtgggg tgcaaaccac gagcgtctac 300
gctgccatga ggggcgcgaa cgccctgggc ccactctgcc tgctgctggc tgccgccacc 360
cagctctcgc ggcagcagtc ccagagaga cctgttttca catgtggtgg cattcttact 420
ggagagtctg gatattattg cagtgaaggt tttcctggag tgtaccctcc aaatagcaaa 480
tgtacttgga aaatcacagt tcccgaagga aaagtagtcg ttctcaattt ccgattcata 540

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gacctcgaga gtgacaacct gtgccgctat gactttgtgg atgtgtacaa tggccatgcc 600
aatggccagc gcattggccg cttctgtggc actttccggc ctggagccct tgtgtccagt 660
ggcaacaaga tgatggtgca gatgatttct gatgccaaca cagctggcaa tggcttcatg 720
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gtgttatttg ttccaccttc aagcctttgc cctgaggtgt tacaatcttg tcttgcgttt 1980
tctaaatcaa tgcttaataa aatattttta aaggaaaaaa aaaaaa 2026

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<210> 104
<211> 415
<212> PRT
<213> Homo sapiens

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<400> 104
Met Arg Gly Ala Asn Ala Trp Ala Pro Leu Cys Leu Leu Leu Ala Ala
 1             5             10             15

Ala Thr Gln Leu Ser Arg Gln Gln Ser Pro Glu Arg Pro Val Phe Thr
          20             25             30

Cys Gly Gly Ile Leu Thr Gly Glu Ser Gly Phe Ile Gly Ser Glu Gly
 35             40             45

Phe Pro Gly Val Tyr Pro Pro Asn Ser Lys Cys Thr Trp Lys Ile Thr
 50             55             60

Val Pro Glu Gly Lys Val Val Val Leu Asn Phe Arg Phe Ile Asp Leu
 65             70             75             80

Glu Ser Asp Asn Leu Cys Arg Tyr Asp Phe Val Asp Val Tyr Asn Gly
          85             90             95

His Ala Asn Gly Gln Arg Ile Gly Arg Phe Cys Gly Thr Phe Arg Pro
          100            105            110

Gly Ala Leu Val Ser Ser Gly Asn Lys Met Met Val Gln Met Ile Ser
          115            120            125

Asp Ala Asn Thr Ala Gly Asn Gly Phe Met Ala Met Phe Ser Ala Ala

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| 130 | 135 | 140 |
|--|-----|-----|
| Glu Pro Asn Glu Arg Gly Asp Gln Tyr Cys Gly Gly Leu Leu Asp Arg 145 150 155 160 | | |
| Pro Ser Gly Ser Phe Lys Thr Pro Asn Trp Pro Asp Arg Asp Tyr Pro 165 170 175 | | |
| Ala Gly Val Thr Cys Val Trp His Ile Val Ala Pro Lys Asn Gln Leu 180 185 190 | | |
| Ile Glu Leu Lys Phe Glu Lys Phe Asp Val Glu Arg Asp Asn Tyr Cys 195 200 205 | | |
| Arg Tyr Asp Tyr Val Ala Val Phe Asn Gly Gly Glu Val Asn Asp Ala 210 215 220 | | |
| Arg Arg Ile Gly Lys Tyr Cys Gly Asp Ser Pro Pro Ala Pro Ile Val 225 230 235 240 | | |
| Ser Glu Arg Asn Glu Leu Leu Ile Gln Phe Leu Ser Asp Leu Ser Leu 245 250 255 | | |
| Thr Ala Asp Gly Phe Ile Gly His Tyr Ile Phe Arg Pro Lys Lys Leu 260 265 270 | | |
| Pro Thr Thr Thr Glu Gln Pro Val Thr Thr Thr Phe Pro Val Thr Thr 275 280 285 | | |
| Gly Leu Lys Pro Thr Val Ala Leu Cys Gln Gln Lys Cys Arg Arg Thr 290 295 300 | | |
| Gly Thr Leu Glu Gly Asn Tyr Cys Ser Ser Asp Phe Val Leu Ala Gly 305 310 315 320 | | |
| Thr Val Ile Thr Thr Ile Thr Arg Asp Gly Ser Leu His Ala Thr Val 325 330 335 | | |
| Ser Ile Ile Asn Ile Tyr Lys Glu Gly Asn Leu Ala Ile Gln Gln Ala 340 345 350 | | |
| Gly Lys Asn Met Ser Ala Arg Leu Thr Val Val Cys Lys Gln Cys Pro 355 360 365 | | |
| Leu Leu Arg Arg Gly Leu Asn Tyr Ile Ile Met Gly Gln Val Gly Glu 370 375 380 | | |
| Asp Gly Arg Gly Lys Ile Met Pro Asn Ser Phe Ile Met Met Phe Lys 385 390 395 400 | | |
| Thr Lys Asn Gln Lys Leu Leu Asp Ala Leu Lys Asn Lys Gln Cys 405 410 415 | | |

<210> 105

<211> 22

<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 105
ccgattcata gacctcgaga gt 22

<210> 106
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 106
gtcaaggagt cctccacaat ac 22

<210> 107
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 107
gtgtacaatg gccatgccaa tggccagcgc attggccgct tctgt 45

<210> 108
<211> 1838
<212> DNA
<213> Homo sapiens

<400> 108
cggacgcgtg ggcggacgcg tgggcggccc acggcgcccg cgggctgggg cggtcgcttc 60
ttccttctcc gtggcctacg aggggtcccca gcctgggtaa agatggcccc atggcccccg 120
aagggcctag tcccagctgt gctctggggc ctcagcctct tcctcaacct cccaggacct 180
atctggctcc agccctctcc acctccccag tcttctcccc cgcctcagcc ccatccgtgt 240
catacctgcc ggggactggt tgacagcttt aacaagggcc tggagagaac catccgggac 300
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gagacccgcc tggtagaggt gctggagggt gtgtgcagca agtcagactt cgagtgccac 420
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gccccggacc tcttccagtg gctgtgctca gattccctga agctctgctg ccccgaggag 540
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gatgagtgtg agacagagggt gtgtccggga gagaacaagc agtgtgaaaa caccgagggc 1080
ggttatcgct gcatctgtgc cgaggggtac aagcagatgg aaggcatctg tgtgaaggag 1140

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cagatcccag agtcagcagg cttcttctca gagatgacag aagacgagtt ggtggtgctg 1200
cagcagatgt tctttggcat catcatctgt gcactggcca cgctggctgc taagggcgac 1260
ttggtgttca ccgccatctt cattggggct gtggcgggcca tgactggcta ctggttgtca 1320
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acaggggtgg ggccatcaca gctccctcct gccagctgca tgctgccagt tcctgttctg 1740
tgttcaccac atccccacac cccattgcca cttatttatt catctcagga aataaagaaa 1800
ggtcttgga agttaaaaaa aaaaaaaaaa aaaaaaaaa 1838

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<210> 109
 <211> 420
 <212> PRT
 <213> Homo sapiens

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<400> 109
Met Ala Pro Trp Pro Pro Lys Gly Leu Val Pro Ala Val Leu Trp Gly
  1              5              10              15

Leu Ser Leu Phe Leu Asn Leu Pro Gly Pro Ile Trp Leu Gln Pro Ser
      20              25              30

Pro Pro Pro Gln Ser Ser Pro Pro Pro Gln Pro His Pro Cys His Thr
      35              40              45

Cys Arg Gly Leu Val Asp Ser Phe Asn Lys Gly Leu Glu Arg Thr Ile
      50              55              60

Arg Asp Asn Phe Gly Gly Gly Asn Thr Ala Trp Glu Glu Glu Asn Leu
      65              70              75              80

Ser Lys Tyr Lys Asp Ser Glu Thr Arg Leu Val Glu Val Leu Glu Gly
      85              90              95

Val Cys Ser Lys Ser Asp Phe Glu Cys His Arg Leu Leu Glu Leu Ser
      100             105             110

Glu Glu Leu Val Glu Ser Trp Trp Phe His Lys Gln Gln Glu Ala Pro
      115             120             125

Asp Leu Phe Gln Trp Leu Cys Ser Asp Ser Leu Lys Leu Cys Cys Pro
      130             135             140

Ala Gly Thr Phe Gly Pro Ser Cys Leu Pro Cys Pro Gly Gly Thr Glu
      145             150             155             160

Arg Pro Cys Gly Gly Tyr Gly Gln Cys Glu Gly Glu Gly Thr Arg Gly
      165             170             175

Gly Ser Gly His Cys Asp Cys Gln Ala Gly Tyr Gly Gly Glu Ala Cys
      180             185             190

Gly Gln Cys Gly Leu Gly Tyr Phe Glu Ala Glu Arg Asn Ala Ser His
      195             200             205

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Leu Val Cys Ser Ala Cys Phe Gly Pro Cys Ala Arg Cys Ser Gly Pro
 210 215 220
 Glu Glu Ser Asn Cys Leu Gln Cys Lys Lys Gly Trp Ala Leu His His
 225 230 235 240
 Leu Lys Cys Val Asp Ile Asp Glu Cys Gly Thr Glu Gly Ala Asn Cys
 245 250 255
 Gly Ala Asp Gln Phe Cys Val Asn Thr Glu Gly Ser Tyr Glu Cys Arg
 260 265 270
 Asp Cys Ala Lys Ala Cys Leu Gly Cys Met Gly Ala Gly Pro Gly Arg
 275 280 285
 Cys Lys Lys Cys Ser Pro Gly Tyr Gln Gln Val Gly Ser Lys Cys Leu
 290 295 300
 Asp Val Asp Glu Cys Glu Thr Glu Val Cys Pro Gly Glu Asn Lys Gln
 305 310 315 320
 Cys Glu Asn Thr Glu Gly Gly Tyr Arg Cys Ile Cys Ala Glu Gly Tyr
 325 330 335
 Lys Gln Met Glu Gly Ile Cys Val Lys Glu Gln Ile Pro Glu Ser Ala
 340 345 350
 Gly Phe Phe Ser Glu Met Thr Glu Asp Glu Leu Val Val Leu Gln Gln
 355 360 365
 Met Phe Phe Gly Ile Ile Ile Cys Ala Leu Ala Thr Leu Ala Ala Lys
 370 375 380
 Gly Asp Leu Val Phe Thr Ala Ile Phe Ile Gly Ala Val Ala Ala Met
 385 390 395 400
 Thr Gly Tyr Trp Leu Ser Glu Arg Ser Asp Arg Val Leu Glu Gly Phe
 405 410 415
 Ile Lys Gly Arg
 420

<210> 110

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 110

cctggctatc agcaggtggg ctccaagtgt ctcgatgtgg atgagtgtga

50

<210> 111

<211> 22

<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 111
attctgcgtg aacactgagg gc

22

<210> 112
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 112
atctgcttgt agccctcggc ac

22

<210> 113
<211> 1616
<212> DNA
<213> Homo sapiens

<220>

<221> modified_base
<222> (1461)..(1461)
<223> a, t, c or g

<400> 113
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cggggccgccc ctgaccgggg agcagctcct gggcagcctg ctgcggcagc tgcagctcaa 180
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cagccagagc ttccgagagg tggccggcag gttcctggcg ttggaggcca gcacacacct 360
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cgacgtgacc gaggccgtga acttctggca gcagctgagc cggccccggc agccgtgctg 660
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acctgagggc agaaagccca ntgtgtcatt gtttacttgt cctgtcactg gatctgggct 1500
aaagtcctcc accaccactc tggacctaag acctgggggtt aagtgtgggt tgtgcatccc 1560
caatccagat aataaagact ttgtaaaaca tgaataaaac acattttatt ctaaaa 1616

<210> 114
<211> 366
<212> PRT
<213> Homo sapiens

<400> 114
Met Gln Pro Leu Trp Leu Cys Trp Ala Leu Trp Val Leu Pro Leu Ala
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Ser Pro Gly Ala Ala Leu Thr Gly Glu Gln Leu Leu Gly Ser Leu Leu
20 25 30
Arg Gln Leu Gln Leu Lys Glu Val Pro Thr Leu Asp Arg Ala Asp Met
35 40 45
Glu Glu Leu Val Ile Pro Thr His Val Arg Ala Gln Tyr Val Ala Leu
50 55 60
Leu Gln Arg Ser His Gly Asp Arg Ser Arg Gly Lys Arg Phe Ser Gln
65 70 75 80
Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Leu Glu Ala Ser Thr
85 90 95
His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro Asn Ser Glu
100 105 110
Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Pro Val Pro Lys Ala
115 120 125
Ala Leu His Arg His Gly Arg Leu Ser Pro Arg Ser Ala Arg Ala Arg
130 135 140
Val Thr Val Glu Trp Leu Arg Val Arg Asp Asp Gly Ser Asn Arg Thr
145 150 155 160
Ser Leu Ile Asp Ser Arg Leu Val Ser Val His Glu Ser Gly Trp Lys
165 170 175
Ala Phe Asp Val Thr Glu Ala Val Asn Phe Trp Gln Gln Leu Ser Arg
180 185 190
Pro Arg Gln Pro Leu Leu Leu Gln Val Ser Val Gln Arg Glu His Leu
195 200 205
Gly Pro Leu Ala Ser Gly Ala His Lys Leu Val Arg Phe Ala Ser Gln
210 215 220
Gly Ala Pro Ala Gly Leu Gly Glu Pro Gln Leu Glu Leu His Thr Leu
225 230 235 240
Asp Leu Gly Asp Tyr Gly Ala Gln Gly Asp Cys Asp Pro Glu Ala Pro
245 250 255

Met Thr Glu Gly Thr Arg Cys Cys Arg Gln Glu Met Tyr Ile Asp Leu
260 265 270

Gln Gly Met Lys Trp Ala Glu Asn Trp Val Leu Glu Pro Pro Gly Phe
275 280 285

Leu Ala Tyr Glu Cys Val Gly Thr Cys Arg Gln Pro Pro Glu Ala Leu
290 295 300

Ala Phe Lys Trp Pro Phe Leu Gly Pro Arg Gln Cys Ile Ala Ser Glu
305 310 315 320

Thr Asp Ser Leu Pro Met Ile Val Ser Ile Lys Glu Gly Gly Arg Thr
325 330 335

Arg Pro Gln Val Val Ser Leu Pro Asn Met Arg Val Gln Lys Cys Ser
340 345 350

Cys Ala Ser Asp Gly Ala Leu Val Pro Arg Arg Leu Gln Pro
355 360 365

<210> 115

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 115

aggactgcca taacttgct g

21

<210> 116

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 116

ataggagttg aagcagcgct gc

22

<210> 117

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 117

tgtgtggaca tagacgagtg ccgctaccgc tactgccagc accgc

45

<210> 118
 <211> 1857
 <212> DNA
 <213> Homo sapiens

<400> 118
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 aaggcgcaag tgcgagaggaa actgttgtgc ctcttcatat tggcgatcct gttgtgctcc 120
 ctggcattgg gcagtgttac agtgcactct tctgaacctg aagtcagaat tcctgagaat 180
 aatcctgtga agttgtcctg tgcctactcg ggcttttctt ctccccgtgt ggagtgggag 240
 tttgaccaag gagacaccac cagactcggt tgctataata acaagatcac agcttcctat 300
 gaggaccggg tgaccttctt gccaaactggt atcaccttca agtccgtgac acgggaagac 360
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 agtgcccga gtgaaggaga attcaaacag acctcgtcac tcctgggtgtg agcctggctg 960
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<210> 119
 <211> 299
 <212> PRT
 <213> Homo sapiens

<400> 119
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 Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu
 35 40 45
 Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe
 50 55 60

Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr
 65 70 75 80
 Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe
 85 90 95
 Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser
 100 105 110
 Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val
 115 120 125
 Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr
 130 135 140
 Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro
 145 150 155 160
 Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn
 165 170 175
 Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro
 180 185 190
 Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly
 195 200 205
 Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser
 210 215 220
 Asn Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val
 225 230 235 240
 Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly
 245 250 255
 Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys Lys Gly
 260 265 270
 Thr Ser Ser Lys Lys Val Ile Tyr Ser Gln Pro Ser Ala Arg Ser Glu
 275 280 285
 Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val
 290 295

<210> 120

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 120

tcgcggagct gtgttctgtt tccc

<210> 121
 <211> 50
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 121
 tgatcgcgat ggggacaaag gcgcaagctc gagaggaaac tgttgtgcct 50

 <210> 122
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 122
 acacctgggtt caaagatggg 20

 <210> 123
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 123
 taggaagagt tgctgaaggc acgg 24

 <210> 124
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 124
 ttgccttact caggtgctac 20

 <210> 125
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 125
actcagcagt ggtaggaaag

20

<210> 126
<211> 1210
<212> DNA
<213> Homo sapiens

<400> 126
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gtcagggctc gtgccacccc accaagttcc agtgccgcac cagtggctta tgcgtgcccc 240
tcacctggcg ctgcgacagg gacttggact gcagcgatgg cagcgatgag gaggagtgca 300
ggattgagcc atgtaccag aaagggcaat gccaccgcc ccctggcctc ccctgcccct 360
gcaccggcgt cagtgactgc tctgggggaa ctgacaagaa actgcgcaac tgcagccgcc 420
tggcctgcct agcaggcgag ctccgttgca cgctgagcga tgactgcatt ccactcacgt 480
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<210> 127
<211> 282
<212> PRT
<213> Homo sapiens

<400> 127
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35 40 45
Pro Ser Ser Gly Ser Cys Pro Pro Thr Lys Phe Gln Cys Arg Thr Ser
50 55 60
Gly Leu Cys Val Pro Leu Thr Trp Arg Cys Asp Arg Asp Leu Asp Cys
65 70 75 80
Ser Asp Gly Ser Asp Glu Glu Glu Cys Arg Ile Glu Pro Cys Thr Gln
85 90 95
Lys Gly Gln Cys Pro Pro Pro Pro Gly Leu Pro Cys Pro Cys Thr Gly

| 100 | 105 | 110 |
|---|-----|-----|
| Val Ser Asp Cys Ser Gly Gly Thr Asp Lys Lys Leu Arg Asn Cys Ser | | |
| 115 | 120 | 125 |
| Arg Leu Ala Cys Leu Ala Gly Glu Leu Arg Cys Thr Leu Ser Asp Asp | | |
| 130 | 135 | 140 |
| Cys Ile Pro Leu Thr Trp Arg Cys Asp Gly His Pro Asp Cys Pro Asp | | |
| 145 | 150 | 155 |
| Ser Ser Asp Glu Leu Gly Cys Gly Thr Asn Glu Ile Leu Pro Glu Gly | | |
| 165 | 170 | 175 |
| Asp Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val Thr Ser | | |
| 180 | 185 | 190 |
| Leu Arg Asn Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val | | |
| 195 | 200 | 205 |
| Pro Ser Val Gly Asn Ala Thr Ser Ser Ser Ala Gly Asp Gln Ser Gly | | |
| 210 | 215 | 220 |
| Ser Pro Thr Ala Tyr Gly Val Ile Ala Ala Ala Ala Val Leu Ser Ala | | |
| 225 | 230 | 235 |
| Ser Leu Val Thr Ala Thr Leu Leu Leu Leu Ser Trp Leu Arg Ala Gln | | |
| 245 | 250 | 255 |
| Glu Arg Leu Arg Pro Leu Gly Leu Leu Val Ala Met Lys Glu Ser Leu | | |
| 260 | 265 | 270 |
| Leu Leu Ser Glu Gln Lys Thr Ser Leu Pro | | |
| 275 | 280 | |

<210> 128

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 128

aagttccagt gccgcaccag tggc

24

<210> 129

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 129

ttggttccac agccgagctc gtcg

24

<210> 130

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 130

gaggaggagt gcaggattga gccatgtacc cagaaagggc aatgccacc

50

<210> 131

<211> 1843

<212> DNA

<213> Homo sapiens

<220>

<221> modified_base

<222> (1837)..(1837)

<223> a, t, c or g

<400> 131

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cagactcttg caagctggat gccctctgtg gatgaaagat gtatcatgga atgaaccoga 180
gcaatggaga tggatttcta gagcagcagc agcagcagca gcaacctcag tccccccaga 240
gactcttggc cgtgatcctg tggtttcagc tggcgtctgt cttcggccct gcacagctca 300
cgggcggggt cgatgacctt caagtgtgtg ctgaccccg cttcccgag aatggcttca 360
ggacccccag cggagggggt ttctttgaag gctctgtagc ccgatttcac tgccaagacg 420
gattcaagct gaagggcgct acaaagagac tgtgtttgaa gcattttaat ggaaccctag 480
gctggatccc aagtataat tccatctgtg tgcaagaaga ttgccgtatc cctcaaactc 540
aagatgctga gattcataac aagacatata gacatggaga gaagctaata atcacttgct 600
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gtttcttctt gacacagact gattaaaaat taaaagnaaa aaa 1843
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<210> 132
 <211> 490
 <212> PRT
 <213> Homo sapiens

<400> 132

| | | | | | | | | | | | | | | | |
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| Met | Tyr | His | Gly | Met | Asn | Pro | Ser | Asn | Gly | Asp | Gly | Phe | Leu | Glu | Gln |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Gln | Gln | Gln | Gln | Gln | Gln | Pro | Gln | Ser | Pro | Gln | Arg | Leu | Leu | Ala | Val |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Ile | Leu | Trp | Phe | Gln | Leu | Ala | Leu | Cys | Phe | Gly | Pro | Ala | Gln | Leu | Thr |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| Gly | Gly | Phe | Asp | Asp | Leu | Gln | Val | Cys | Ala | Asp | Pro | Gly | Ile | Pro | Glu |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| Asn | Gly | Phe | Arg | Thr | Pro | Ser | Gly | Gly | Val | Phe | Phe | Glu | Gly | Ser | Val |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| Ala | Arg | Phe | His | Cys | Gln | Asp | Gly | Phe | Lys | Leu | Lys | Gly | Ala | Thr | Lys |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| Arg | Leu | Cys | Leu | Lys | His | Phe | Asn | Gly | Thr | Leu | Gly | Trp | Ile | Pro | Ser |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Asp | Asn | Ser | Ile | Cys | Val | Gln | Glu | Asp | Cys | Arg | Ile | Pro | Gln | Ile | Glu |
| | | 115 | | | | | 120 | | | | | 125 | | | |
| Asp | Ala | Glu | Ile | His | Asn | Lys | Thr | Tyr | Arg | His | Gly | Glu | Lys | Leu | Ile |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| Ile | Thr | Cys | His | Glu | Gly | Phe | Lys | Ile | Arg | Tyr | Pro | Asp | Leu | His | Asn |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Met | Val | Ser | Leu | Cys | Arg | Asp | Asp | Gly | Thr | Trp | Asn | Asn | Leu | Pro | Ile |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Cys | Gln | Gly | Cys | Leu | Arg | Pro | Leu | Ala | Ser | Ser | Asn | Gly | Tyr | Val | Asn |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Ile | Ser | Glu | Leu | Gln | Thr | Ser | Phe | Pro | Val | Gly | Thr | Val | Ile | Ser | Tyr |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Arg | Cys | Phe | Pro | Gly | Phe | Lys | Leu | Asp | Gly | Ser | Ala | Tyr | Leu | Glu | Cys |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Leu | Gln | Asn | Leu | Ile | Trp | Ser | Ser | Ser | Pro | Pro | Arg | Cys | Leu | Ala | Leu |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Glu | Ala | Gln | Val | Cys | Pro | Leu | Pro | Pro | Met | Val | Ser | His | Gly | Asp | Phe |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Val | Cys | His | Pro | Arg | Pro | Cys | Glu | Arg | Tyr | Asn | His | Gly | Thr | Val | Val |
| | | | 260 | | | | | 265 | | | | | 270 | | |

Glu Phe Tyr Cys Asp Pro Gly Tyr Ser Leu Thr Ser Asp Tyr Lys Tyr
 275 280 285
 Ile Thr Cys Gln Tyr Gly Glu Trp Phe Pro Ser Tyr Gln Val Tyr Cys
 290 295 300
 Ile Lys Ser Glu Gln Thr Trp Pro Ser Thr His Glu Thr Leu Leu Thr
 305 310 315 320
 Thr Trp Lys Ile Val Ala Phe Thr Ala Thr Ser Val Leu Leu Val Leu
 325 330 335
 Leu Leu Val Ile Leu Ala Arg Met Phe Gln Thr Lys Phe Lys Ala His
 340 345 350
 Phe Pro Pro Arg Gly Pro Pro Arg Ser Ser Ser Ser Asp Pro Asp Phe
 355 360 365
 Val Val Val Asp Gly Val Pro Val Met Leu Pro Ser Tyr Asp Glu Ala
 370 375 380
 Val Ser Gly Gly Leu Ser Ala Leu Gly Pro Gly Tyr Met Ala Ser Val
 385 390 395 400
 Gly Gln Gly Cys Pro Leu Pro Val Asp Asp Gln Ser Pro Pro Ala Tyr
 405 410 415
 Pro Gly Ser Gly Asp Thr Asp Thr Gly Pro Gly Glu Ser Glu Thr Cys
 420 425 430
 Asp Ser Val Ser Gly Ser Ser Glu Leu Leu Gln Ser Leu Tyr Ser Pro
 435 440 445
 Pro Arg Cys Gln Glu Ser Thr His Pro Ala Ser Asp Asn Pro Asp Ile
 450 455 460
 Ile Ala Ser Thr Ala Glu Glu Val Ala Ser Thr Ser Pro Gly Ile His
 465 470 475 480
 His Ala His Trp Val Leu Phe Leu Arg Asn
 485 490

<210> 133

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 133

atctctatc gctgctttcc cgg

23

<210> 134

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 134

agccaggatc gcagtaaaac tcc

23

<210> 135

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 135

atttaaactt gatgggtctg cgtatcttga gtgcttacaa aaccttatct

50

<210> 136

<211> 1815

<212> DNA

<213> Homo sapiens

<400> 136

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agcaggaaaa aaaaa

1815

<210> 137

<211> 382

<212> PRT

<213> Homo sapiens

<400> 137

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Val Gly Leu Arg Ala Ala Thr Gly Arg Leu Leu Ser Ala Ser Asp Leu
20 25 30

Asp Leu Arg Gly Gly Gln Pro Val Cys Arg Gly Gly Thr Gln Arg Pro
35 40 45

Cys Tyr Lys Val Ile Tyr Phe His Asp Thr Ser Arg Arg Leu Asn Phe
50 55 60

Glu Glu Ala Lys Glu Ala Cys Arg Arg Asp Gly Gly Gln Leu Val Ser
65 70 75 80

Ile Glu Ser Glu Asp Glu Gln Lys Leu Ile Glu Lys Phe Ile Glu Asn
85 90 95

Leu Leu Pro Ser Asp Gly Asp Phe Trp Ile Gly Leu Arg Arg Arg Glu
100 105 110

Glu Lys Gln Ser Asn Ser Thr Ala Cys Gln Asp Leu Tyr Ala Trp Thr
115 120 125

Asp Gly Ser Ile Ser Gln Phe Arg Asn Trp Tyr Val Asp Glu Pro Ser
130 135 140

Cys Gly Ser Glu Val Cys Val Val Met Tyr His Gln Pro Ser Ala Pro
145 150 155 160

Ala Gly Ile Gly Gly Pro Tyr Met Phe Gln Trp Asn Asp Asp Arg Cys
165 170 175

Asn Met Lys Asn Asn Phe Ile Cys Lys Tyr Ser Asp Glu Lys Pro Ala
180 185 190

Val Pro Ser Arg Glu Ala Glu Gly Glu Glu Thr Glu Leu Thr Thr Pro
195 200 205

Val Leu Pro Glu Glu Thr Gln Glu Glu Asp Ala Lys Lys Thr Phe Lys
210 215 220

Glu Ser Arg Glu Ala Ala Leu Asn Leu Ala Tyr Ile Leu Ile Pro Ser
225 230 235 240

Ile Pro Leu Leu Leu Leu Leu Val Val Thr Thr Val Val Cys Trp Val
245 250 255

Trp Ile Cys Arg Lys Arg Lys Arg Glu Gln Pro Asp Pro Ser Thr Lys

| 260 | | | | | | | | | | 265 | | | | | 270 | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|--|--|--|--|--|--|--|--|--|--|--|--|--|--|
| Lys | Gln | His | Thr | Ile | Trp | Pro | Ser | Pro | His | Gln | Gly | Asn | Ser | Pro | Asp | | | | | | | | | | | | | | | |
| | | 275 | | | | | 280 | | | | | 285 | | | | | | | | | | | | | | | | | | |
| Leu | Glu | Val | Tyr | Asn | Val | Ile | Arg | Lys | Gln | Ser | Glu | Ala | Asp | Leu | Ala | | | | | | | | | | | | | | | |
| | 290 | | | | | 295 | | | | | 300 | | | | | | | | | | | | | | | | | | | |
| Glu | Thr | Arg | Pro | Asp | Leu | Lys | Asn | Ile | Ser | Phe | Arg | Val | Cys | Ser | Gly | | | | | | | | | | | | | | | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | | | | | | | | | | | | | | | |
| Glu | Ala | Thr | Pro | Asp | Asp | Met | Ser | Cys | Asp | Tyr | Asp | Asn | Met | Ala | Val | | | | | | | | | | | | | | | |
| | | | | 325 | | | | | 330 | | | | | 335 | | | | | | | | | | | | | | | | |
| Asn | Pro | Ser | Glu | Ser | Gly | Phe | Val | Thr | Leu | Val | Ser | Val | Glu | Ser | Gly | | | | | | | | | | | | | | | |
| | | | 340 | | | | | 345 | | | | | 350 | | | | | | | | | | | | | | | | | |
| Phe | Val | Thr | Asn | Asp | Ile | Tyr | Glu | Phe | Ser | Pro | Asp | Gln | Met | Gly | Arg | | | | | | | | | | | | | | | |
| | 355 | | | | | 360 | | | | | | 365 | | | | | | | | | | | | | | | | | | |
| Ser | Lys | Glu | Ser | Gly | Trp | Val | Glu | Asn | Glu | Ile | Tyr | Gly | Tyr | | | | | | | | | | | | | | | | | |
| | 370 | | | | | 375 | | | | | 380 | | | | | | | | | | | | | | | | | | | |

<210> 138
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 138
 gttcattgaa aacctcttgc catctgatgg tgacttctgg attgggctca 50

<210> 139
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 139
 aagccaaaga agcctgcagg aggg 24

<210> 140
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 140

cagtccaagc ataaaggtcc tggc

24

<210> 141
<211> 1514
<212> DNA
<213> Homo sapiens

<400> 141
ggggtctccc tcagggccgg gaggcacagc ggtccctgct tgctgaaggg ctggatgtac 60
gcatccgcag gttcccgagg acttgggggc gcccgctgag ccccggcgcc cgcagaagac 120
ttgtgtttgc ctccctgcagc ctcaaccggg agggcagcga gggcctacca ccatgatcac 180
tggtgtgttc agcatgcgct tgtggacccc agtgggcgtc ctgacctcgc tggcgactcg 240
cctgcaccag cggcggttgg ccctggccga gctgcaggag gccgatggcc agtgtccggt 300
cgaccgcagc ctgctgaagt tgaaaatggt gcaggtcgtg tttcgacacg gggctcggag 360
tcctctcaag ccgctcccg c tgaggagca ggtagagtgg aacccccagc tattagaggt 420
cccaccccaa actcagtttg attacacagt caccaatcta gctggtggtc cgaaaccata 480
ttctccttac gactctcaat accatgagac caccctgaag gggggcatgt ttgctgggca 540
gctgaccaag gtgggcattgc agcaaatggt tgccttgga gagagactga ggaagaacta 600
tgttgaagac attccctttc tttcaccaac cttcaaccga caggaggtct ttattcggtc 660
cactaacatt tttcggaatc tggagtccac ccgttgttg ctggctgggc ttttccagtg 720
tcagaaagaa ggacccatca tcatccacac tgatgaagca gattcagaag tcttgtatcc 780
caactaccaa agctgctgga gcctgaggca gagaaccaga ggccggaggc agactgcctc 840
tttacagcca ggaatctcag aggatttgaa aaaggatgaag gacaggatgg gcattgacag 900
tagtgataaa gtggacttct tcatcctcct ggacaacgtg gctgccgagc aggcacacaa 960
cctcccaagc tgcccatgc tgaagagatt tgcacggatg atcgaacaga gagctgtgga 1020
cacatccttg tacatactgc ccaaggaaga cagggaagt cttcagatgg cagtagggcc 1080
attcctccac atcctagaga gcaacctgct gaaagccatg gactctgcca ctgcccccca 1140
caagatcaga aagctgtatc tctatgcggc tcatgatgtg accttcatac cgctcttaat 1200
gaccctgggg atttttgacc acaaatggc accgtttgct gttgacctga ccatggaact 1260
ttaccagcac ctggaatcta aggagtgggt tgtgcagctc tattaccacg ggaaggagca 1320
gggtgccgaga ggttgccctg atgggctctg cccgctggac atgttcttga atgccatgtc 1380
agtttatacc ttaagcccag aaaaatacca tgcactctgc tctcaaaactc aggtgatgga 1440
agtttgaaat gaagagtaac tgatttataa aagcaggatg tgttgatttt aaaataaagt 1500
gcctttatac aatg 1514

<210> 142
<211> 428
<212> PRT
<213> Homo sapiens

<400> 142
Met Ile Thr Gly Val Phe Ser Met Arg Leu Trp Thr Pro Val Gly Val
1 5 10 15
Leu Thr Ser Leu Ala Tyr Cys Leu His Gln Arg Arg Val Ala Leu Ala
20 25 30
Glu Leu Gln Glu Ala Asp Gly Gln Cys Pro Val Asp Arg Ser Leu Leu
35 40 45
Lys Leu Lys Met Val Gln Val Val Phe Arg His Gly Ala Arg Ser Pro
50 55 60
Leu Lys Pro Leu Pro Leu Glu Glu Gln Val Glu Trp Asn Pro Gln Leu
65 70 75 80
Leu Glu Val Pro Pro Gln Thr Gln Phe Asp Tyr Thr Val Thr Asn Leu

| 85 | | | | | | | | 90 | | | | 95 | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Ala | Gly | Gly | Pro | Lys | Pro | Tyr | Ser | Pro | Tyr | Asp | Ser | Gln | Tyr | His | Glu | |
| | | | 100 | | | | | 105 | | | | | 110 | | | |
| Thr | Thr | Leu | Lys | Gly | Gly | Met | Phe | Ala | Gly | Gln | Leu | Thr | Lys | Val | Gly | |
| | | 115 | | | | | 120 | | | | | 125 | | | | |
| Met | Gln | Gln | Met | Phe | Ala | Leu | Gly | Glu | Arg | Leu | Arg | Lys | Asn | Tyr | Val | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Glu | Asp | Ile | Pro | Phe | Leu | Ser | Pro | Thr | Phe | Asn | Pro | Gln | Glu | Val | Phe | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | |
| Ile | Arg | Ser | Thr | Asn | Ile | Phe | Arg | Asn | Leu | Glu | Ser | Thr | Arg | Cys | Leu | |
| | | | | 165 | | | | | 170 | | | | | 175 | | |
| Leu | Ala | Gly | Leu | Phe | Gln | Cys | Gln | Lys | Glu | Gly | Pro | Ile | Ile | Ile | His | |
| | | | 180 | | | | | | 185 | | | | 190 | | | |
| Thr | Asp | Glu | Ala | Asp | Ser | Glu | Val | Leu | Tyr | Pro | Asn | Tyr | Gln | Ser | Cys | |
| | | 195 | | | | | 200 | | | | | 205 | | | | |
| Trp | Ser | Leu | Arg | Gln | Arg | Thr | Arg | Gly | Arg | Arg | Gln | Thr | Ala | Ser | Leu | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Gln | Pro | Gly | Ile | Ser | Glu | Asp | Leu | Lys | Lys | Val | Lys | Asp | Arg | Met | Gly | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | |
| Ile | Asp | Ser | Ser | Asp | Lys | Val | Asp | Phe | Phe | Ile | Leu | Leu | Asp | Asn | Val | |
| | | | | 245 | | | | | 250 | | | | | 255 | | |
| Ala | Ala | Glu | Gln | Ala | His | Asn | Leu | Pro | Ser | Cys | Pro | Met | Leu | Lys | Arg | |
| | | | 260 | | | | | | 265 | | | | 270 | | | |
| Phe | Ala | Arg | Met | Ile | Glu | Gln | Arg | Ala | Val | Asp | Thr | Ser | Leu | Tyr | Ile | |
| | | 275 | | | | | 280 | | | | | 285 | | | | |
| Leu | Pro | Lys | Glu | Asp | Arg | Glu | Ser | Leu | Gln | Met | Ala | Val | Gly | Pro | Phe | |
| | 290 | | | | | 295 | | | | | 300 | | | | | |
| Leu | His | Ile | Leu | Glu | Ser | Asn | Leu | Leu | Lys | Ala | Met | Asp | Ser | Ala | Thr | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | |
| Ala | Pro | Asp | Lys | Ile | Arg | Lys | Leu | Tyr | Leu | Tyr | Ala | Ala | His | Asp | Val | |
| | | | | 325 | | | | | 330 | | | | | 335 | | |
| Thr | Phe | Ile | Pro | Leu | Leu | Met | Thr | Leu | Gly | Ile | Phe | Asp | His | Lys | Trp | |
| | | | 340 | | | | | | 345 | | | | 350 | | | |
| Pro | Pro | Phe | Ala | Val | Asp | Leu | Thr | Met | Glu | Leu | Tyr | Gln | His | Leu | Glu | |
| | | 355 | | | | | 360 | | | | | 365 | | | | |
| Ser | Lys | Glu | Trp | Phe | Val | Gln | Leu | Tyr | Tyr | His | Gly | Lys | Glu | Gln | Val | |
| | 370 | | | | | 375 | | | | | 380 | | | | | |
| Pro | Arg | Gly | Cys | Pro | Asp | Gly | Leu | Cys | Pro | Leu | Asp | Met | Phe | Leu | Asn | |

| | | | |
|---|---|-----|-----|
| 385 | 390 | 395 | 400 |
| Ala Met Ser Val | Tyr Thr Leu Ser Pro Glu Lys Tyr His Ala Leu Cys | | |
| | 405 | 410 | 415 |
| Ser Gln Thr Gln Val Met Glu Val Gly Asn Glu Glu | | | |
| | 420 | 425 | |

<210> 143
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 143
 ccaactacca aagctgctgg agcc 24

<210> 144
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 144
 gcagctctat taccacggga agga 24

<210> 145
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 145
 tccttcccgt ggtaatagag ctgc 24

<210> 146
 <211> 45
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 146
 ggcagagaac cagaggccgg aggagactgc ctctttacag ccagg 45

<210> 147

<211> 1686
<212> DNA
<213> Homo sapiens

<400> 147
ctcctcttaa cataacttgca gctaaaacta aatattgctg cttggggacc tccttctagc 60
cttaaatttc agctcatcac cttcacctgc cttgggtcatg gctctgctat tctccttgat 120
ccttgccatt tgcaccagac ctggattcct agcgtctcca tctggagtgc ggctgggtggg 180
gggcctccac cgctgtgaag ggcgggtgga ggtggaacag aaaggccagt ggggcaccgt 240
gtgtgatgac ggctgggaca ttaaggacgt ggctgtgttg tgccgggagc tgggctgtgg 300
agctgccagc ggaaccccta gtggtatatt gtatgagcca ccagcagaaa aagagcaaaa 360
ggtcctcatc caatcagtca gttgcacagg aacagaagat acattggctc agtgtgagca 420
agaagaagtt tatgattggt cacatgatga agatgctggg gcatcgtgtg agaaccaga 480
gagctctttc tccccagtc cagaggggtgt caggctggct gacggccctg ggcatcgcaa 540
gggacgcgtg gaagtgaagc accagaacca gtggtatacc gtgtgccaga caggctggag 600
cctccgggcc gcaaaggtgg tgtgccggca gctgggatgt gggagggtg tactgactca 660
aaaacgctgc aacaagcatg cctatggccg aaaacccatc tggctgagcc agatgtcatg 720
ctcaggacga gaagcaaccc ttcaggattg cccttctggg ccttggggga agaacacctg 780
caacatgat gaagacacgt gggtcgaatg tgaagatccc tttgacttga gactagtagg 840
aggagacaac ctctgctctg ggcgactgga ggtgctgcac aagggcgtat ggggctctgt 900
ctgtgatgac aactggggag aaaaggagga ccagggtgta tgcaagcaac tgggctgtgg 960
gaagtccctc tctccctcct tcagagaccg gaaatgctat ggccctgggg ttggccgcat 1020
ctggctggat aatgttcgtt gctcagggga ggagcagtc ctggagcagt gccagcacag 1080
attttggggg tttcacgact gcacccacca ggaagatgtg gctgtcatct gctcagtgtg 1140
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atttactgtc tacatgactg catgggatga acactgatct tcttctgcc ttggactggg 1260
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aggtctagtt ctcaggccat cagacatagt ttggaactac atcaccacct ttcctatgtc 1380
tccacattgc acacagcaga ttcccagcct ccataattgt gtgtatcaac tacttaaata 1440
cattctcaca cacacacaca cacacacaca cacacacaca ccatttgtcc 1500
tgtttctctg aagaactctg acaaaataca gatttttgta ctgaaagaga ttctagagga 1560
acggaatttt aaggataaat tttctgaatt ggttatgggg tttctgaaat tggctctata 1620
atctaattag atataaaaatt ctggttaact tatttacaat aataaagata gcactatgtg 1680
ttcaaa 1686

<210> 148
<211> 347
<212> PRT
<213> Homo sapiens

<400> 148
Met Ala Leu Leu Phe Ser Leu Ile Leu Ala Ile Cys Thr Arg Pro Gly
1 5 10 15
Phe Leu Ala Ser Pro Ser Gly Val Arg Leu Val Gly Gly Leu His Arg
20 25 30
Cys Glu Gly Arg Val Glu Val Glu Gln Lys Gly Gln Trp Gly Thr Val
35 40 45
Cys Asp Asp Gly Trp Asp Ile Lys Asp Val Ala Val Leu Cys Arg Glu
50 55 60
Leu Gly Cys Gly Ala Ala Ser Gly Thr Pro Ser Gly Ile Leu Tyr Glu
65 70 75 80
Pro Pro Ala Glu Lys Glu Gln Lys Val Leu Ile Gln Ser Val Ser Cys

| 85 | | | | | 90 | | | | | 95 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Thr | Gly | Thr | Glu | Asp | Thr | Leu | Ala | Gln | Cys | Glu | Gln | Glu | Glu | Val | Tyr |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Asp | Cys | Ser | His | Asp | Glu | Asp | Ala | Gly | Ala | Ser | Cys | Glu | Asn | Pro | Glu |
| | | 115 | | | | | 120 | | | | | 125 | | | |
| Ser | Ser | Phe | Ser | Pro | Val | Pro | Glu | Gly | Val | Arg | Leu | Ala | Asp | Gly | Pro |
| | | 130 | | | | 135 | | | | | 140 | | | | |
| Gly | His | Cys | Lys | Gly | Arg | Val | Glu | Val | Lys | His | Gln | Asn | Gln | Trp | Tyr |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Thr | Val | Cys | Gln | Thr | Gly | Trp | Ser | Leu | Arg | Ala | Ala | Lys | Val | Val | Cys |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Arg | Gln | Leu | Gly | Cys | Gly | Arg | Ala | Val | Leu | Thr | Gln | Lys | Arg | Cys | Asn |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Lys | His | Ala | Tyr | Gly | Arg | Lys | Pro | Ile | Trp | Leu | Ser | Gln | Met | Ser | Cys |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Ser | Gly | Arg | Glu | Ala | Thr | Leu | Gln | Asp | Cys | Pro | Ser | Gly | Pro | Trp | Gly |
| | | 210 | | | | 215 | | | | | 220 | | | | |
| Lys | Asn | Thr | Cys | Asn | His | Asp | Glu | Asp | Thr | Trp | Val | Glu | Cys | Glu | Asp |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Pro | Phe | Asp | Leu | Arg | Leu | Val | Gly | Gly | Asp | Asn | Leu | Cys | Ser | Gly | Arg |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Leu | Glu | Val | Leu | His | Lys | Gly | Val | Trp | Gly | Ser | Val | Cys | Asp | Asp | Asn |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Trp | Gly | Glu | Lys | Glu | Asp | Gln | Val | Val | Cys | Lys | Gln | Leu | Gly | Cys | Gly |
| | | 275 | | | | 280 | | | | | 285 | | | | |
| Lys | Ser | Leu | Ser | Pro | Ser | Phe | Arg | Asp | Arg | Lys | Cys | Tyr | Gly | Pro | Gly |
| | | 290 | | | | 295 | | | | | 300 | | | | |
| Val | Gly | Arg | Ile | Trp | Leu | Asp | Asn | Val | Arg | Cys | Ser | Gly | Glu | Glu | Gln |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Ser | Leu | Glu | Gln | Cys | Gln | His | Arg | Phe | Trp | Gly | Phe | His | Asp | Cys | Thr |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| His | Gln | Glu | Asp | Val | Ala | Val | Ile | Cys | Ser | Val | | | | | |
| | | | 340 | | | | | 345 | | | | | | | |

<210> 149

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

oligonucleotide probe

<400> 149

ttcagctcat caccttcacc tgcc

24

<210> 150

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 150

ggctcataca aaataccact aggg

24

<210> 151

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 151

gggcctccac cgctgtgaag ggcgggtgga ggtggaacag aaaggccagt

50

<210> 152

<211> 1427

<212> DNA

<213> Homo sapiens

<400> 152

| | | | | | | |
|-------------|------------|------------|-------------|-------------|------------|------|
| actgcactcg | gttctatcga | ttgaattccc | cggggatcct | ctagagatcc | ctcgacctcg | 60 |
| acccacgcgt | ccgcggacgc | gtgggcggac | gcgtgggccg | gctaccagga | agagtctgcc | 120 |
| gaagggtgaag | gccatggact | tcatacctc | cacagccatc | ctgcccctgc | tgttcggtcg | 180 |
| cctgggcgtc | ttcggcctct | tccggctgct | gcagtgggtg | cgcggaag | cctacctgcg | 240 |
| gaatgctgtg | gtggtgatca | caggcgccac | ctcagggtcg | ggcaaagaat | gtgcaaaagt | 300 |
| cttctatgct | gcgggtgcta | aactggtgct | ctgtggccgg | aatggtggg | ccctagaaga | 360 |
| gctcatcaga | gaacttaccg | cttctcatgc | caccaagggtg | cagacacaca | agccttactt | 420 |
| ggtgaccttc | gacctcacag | actctggggc | catagttgca | gcagcagctg | agatcctgca | 480 |
| gtgctttggc | tatgtcgaca | tacttgtcaa | caatgctggg | atcagctacc | gtggtaccat | 540 |
| catggacacc | acagtggatg | tggacaagag | ggtcatggag | acaaactact | ttggcccagt | 600 |
| tgctctaacg | aaagcactcc | tgccctccat | gatcaagagg | aggcaaggcc | acattgtcgc | 660 |
| catcagcagc | atccagggca | agatgagcat | tccttttcga | tcagcatatg | cagcctccaa | 720 |
| gcacgcaacc | caggctttct | ttgactgtct | gcgtgccgag | atggaacagt | atgaaattga | 780 |
| ggtgaccgtc | atcagccccg | gctacatcca | caccaacctc | tctgtaaattg | ccatcaccgc | 840 |
| ggatggatct | aggtatggag | ttatggacac | caccacagcc | cagggccgaa | gccctgtgga | 900 |
| ggtggcccag | gatgttcttg | ctgctgtggg | gaagaagaag | aaagatgtga | tcctggctga | 960 |
| cttactgcct | tccttggtcg | tttatcttcg | aactctggct | cctgggctct | tcttcagcct | 1020 |
| catggcctcc | agggccagaa | aagagcggaa | atccaagaac | tcctagtact | ctgaccagcc | 1080 |
| agggccagg | cagagaagca | gcactcttag | gcttgcttac | tctacaagg | acagttgcat | 1140 |
| ttgttgagac | tttaatggag | atgtgtctca | caagtgggaa | agactgaaga | aacacatctc | 1200 |
| gtgcagatct | gctggcagag | gacaatcaaa | aacgacaaca | agcttcttcc | cagggtgagg | 1260 |
| ggaaacactt | aaggaataaa | tatggagctg | gggtttaaca | ctaaaaacta | gaaataaaca | 1320 |

tctcaaacag taaaaaaaaa aaaaaagggc ggccgcgact ctagagtcga cctgcagaag 1380
 cttggccgcc atggcccaac ttgtttattg cagcttataa tgggttac 1427

<210> 153
 <211> 310
 <212> PRT
 <213> Homo sapiens

<400> 153

| | | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|
| Met | Asp | Phe | Ile | Thr | Ser | Thr | Ala | Ile | Leu | Pro | Leu | Leu | Phe | Gly | Cys | 1 | 5 | 10 | 15 |
| Leu | Gly | Val | Phe | Gly | Leu | Phe | Arg | Leu | Leu | Gln | Trp | Val | Arg | Gly | Lys | 20 | 25 | 30 | |
| Ala | Tyr | Leu | Arg | Asn | Ala | Val | Val | Val | Ile | Thr | Gly | Ala | Thr | Ser | Gly | 35 | 40 | 45 | |
| Leu | Gly | Lys | Glu | Cys | Ala | Lys | Val | Phe | Tyr | Ala | Ala | Gly | Ala | Lys | Leu | 50 | 55 | 60 | |
| Val | Leu | Cys | Gly | Arg | Asn | Gly | Gly | Ala | Leu | Glu | Glu | Leu | Ile | Arg | Glu | 65 | 70 | 75 | |
| Leu | Thr | Ala | Ser | His | Ala | Thr | Lys | Val | Gln | Thr | His | Lys | Pro | Tyr | Leu | 85 | 90 | 95 | |
| Val | Thr | Phe | Asp | Leu | Thr | Asp | Ser | Gly | Ala | Ile | Val | Ala | Ala | Ala | Ala | 100 | 105 | 110 | |
| Glu | Ile | Leu | Gln | Cys | Phe | Gly | Tyr | Val | Asp | Ile | Leu | Val | Asn | Asn | Ala | 115 | 120 | 125 | |
| Gly | Ile | Ser | Tyr | Arg | Gly | Thr | Ile | Met | Asp | Thr | Thr | Val | Asp | Val | Asp | 130 | 135 | 140 | |
| Lys | Arg | Val | Met | Glu | Thr | Asn | Tyr | Phe | Gly | Pro | Val | Ala | Leu | Thr | Lys | 145 | 150 | 155 | |
| Ala | Leu | Leu | Pro | Ser | Met | Ile | Lys | Arg | Arg | Gln | Gly | His | Ile | Val | Ala | 165 | 170 | 175 | |
| Ile | Ser | Ser | Ile | Gln | Gly | Lys | Met | Ser | Ile | Pro | Phe | Arg | Ser | Ala | Tyr | 180 | 185 | 190 | |
| Ala | Ala | Ser | Lys | His | Ala | Thr | Gln | Ala | Phe | Phe | Asp | Cys | Leu | Arg | Ala | 195 | 200 | 205 | |
| Glu | Met | Glu | Gln | Tyr | Glu | Ile | Glu | Val | Thr | Val | Ile | Ser | Pro | Gly | Tyr | 210 | 215 | 220 | |
| Ile | His | Thr | Asn | Leu | Ser | Val | Asn | Ala | Ile | Thr | Ala | Asp | Gly | Ser | Arg | 225 | 230 | 235 | |
| Tyr | Gly | Val | Met | Asp | Thr | Thr | Thr | Ala | Gln | Gly | Arg | Ser | Pro | Val | Glu | 245 | 250 | 255 | |

Val Ala Gln Asp Val Leu Ala Ala Val Gly Lys Lys Lys Lys Asp Val
260 265 270

Ile Leu Ala Asp Leu Leu Pro Ser Leu Ala Val Tyr Leu Arg Thr Leu
275 280 285

Ala Pro Gly Leu Phe Phe Ser Leu Met Ala Ser Arg Ala Arg Lys Glu
290 295 300

Arg Lys Ser Lys Asn Ser
305 310

<210> 154

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 154

ggtgctaaac tgggtgctctg tggc

24

<210> 155

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 155

cagggcaaga tgagcattcc

20

<210> 156

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 156

tcatactgtt ccatctcggc acgc

24

<210> 157

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 157
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<210> 158
<211> 1771
<212> DNA
<213> Homo sapiens

<400> 158
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cctcctgctt ctcccgttac tgatcgtctg ctccctagag tccttcgtga agctttttat 180
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aattgggaga ctgactgcct atgaatttgc taaacttaaa agcaagctgg ttctctggga 300
tataaataag catggactgg aggaaacagc tgccaaatgc aaggggactgg gtgccaaggt 360
tcataccttt gtggtagact gcagcaaccg agaagatatt tacagctctg caaagaaggt 420
gaaggcagaa attggagatg ttagtatttt agtaaataat gctggtgtag tctatacatc 480
agatttgttt gctacacaag atcctcagat tgaaaagact tttgaagtta atgtacttgc 540
acatttctgg actacaaagg catttcttcc tgcaatgacg aagaataacc atggccatat 600
tgtcactgtg gcttcggcag ctggacatgt ctcggtcccc ttcttactgg cttactgttc 660
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gccactctgt ttcttgagag atacctcaca ttccaatgcc aaacatttct gcacagggaa 1560
gctagagggtg gatacacgtg ttgcaagtat aaaagcatca ctgggattta aggagaattg 1620
agagaatgta cccacaaatg gcagcaataa taaatggatc acacttaaaa aaaaaaaaaa 1680
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<210> 159
<211> 300
<212> PRT
<213> Homo sapiens

<400> 159
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Cys Ser Leu Glu Ser Phe Val Lys Leu Phe Ile Pro Lys Arg Arg Lys
20 25 30
Ser Val Thr Gly Glu Ile Val Leu Ile Thr Gly Ala Gly His Gly Ile
35 40 45
Gly Arg Leu Thr Ala Tyr Glu Phe Ala Lys Leu Lys Ser Lys Leu Val
50 55 60

| | | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Trp | Asp | Ile | Asn | Lys | His | Gly | Leu | Glu | Glu | Thr | Ala | Ala | Lys | Cys | 65 | 70 | 75 | 80 |
| Lys | Gly | Leu | Gly | Ala | Lys | Val | His | Thr | Phe | Val | Val | Asp | Cys | Ser | Asn | 85 | 90 | 95 | |
| Arg | Glu | Asp | Ile | Tyr | Ser | Ser | Ala | Lys | Lys | Val | Lys | Ala | Glu | Ile | Gly | 100 | 105 | 110 | |
| Asp | Val | Ser | Ile | Leu | Val | Asn | Asn | Ala | Gly | Val | Val | Tyr | Thr | Ser | Asp | 115 | 120 | 125 | |
| Leu | Phe | Ala | Thr | Gln | Asp | Pro | Gln | Ile | Glu | Lys | Thr | Phe | Glu | Val | Asn | 130 | 135 | 140 | |
| Val | Leu | Ala | His | Phe | Trp | Thr | Thr | Lys | Ala | Phe | Leu | Pro | Ala | Met | Thr | 145 | 150 | 155 | 160 |
| Lys | Asn | Asn | His | Gly | His | Ile | Val | Thr | Val | Ala | Ser | Ala | Ala | Gly | His | 165 | 170 | 175 | |
| Val | Ser | Val | Pro | Phe | Leu | Leu | Ala | Tyr | Cys | Ser | Ser | Lys | Phe | Ala | Ala | 180 | 185 | 190 | |
| Val | Gly | Phe | His | Lys | Thr | Leu | Thr | Asp | Glu | Leu | Ala | Ala | Leu | Gln | Ile | 195 | 200 | 205 | |
| Thr | Gly | Val | Lys | Thr | Thr | Cys | Leu | Cys | Pro | Asn | Phe | Val | Asn | Thr | Gly | 210 | 215 | 220 | |
| Phe | Ile | Lys | Asn | Pro | Ser | Thr | Ser | Leu | Gly | Pro | Thr | Leu | Glu | Pro | Glu | 225 | 230 | 235 | 240 |
| Glu | Val | Val | Asn | Arg | Leu | Met | His | Gly | Ile | Leu | Thr | Glu | Gln | Lys | Met | 245 | 250 | 255 | |
| Ile | Phe | Ile | Pro | Ser | Ser | Ile | Ala | Phe | Leu | Thr | Thr | Leu | Glu | Arg | Ile | 260 | 265 | 270 | |
| Leu | Pro | Glu | Arg | Phe | Leu | Ala | Val | Leu | Lys | Arg | Lys | Ile | Ser | Val | Lys | 275 | 280 | 285 | |
| Phe | Asp | Ala | Val | Ile | Gly | Tyr | Lys | Met | Lys | Ala | Gln | | | | | 290 | 295 | 300 | |

<210> 160

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 160

ggtgaaggca gaaattggag atg

<210> 161
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 161
atcccatgca tcagcctggt tacc 24

<210> 162
<211> 48
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 162
gctggtgtag tctatacatc agatttggtt gctacacaag atcctcag 48

<210> 163
<211> 2076
<212> DNA
<213> Homo sapiens

<400> 163
cccacgcgtc cgcggacgcg tgggtcgact agttctagat cgcgagcggc cgcccgcggc 60
tcaggaggga gcaccgactg cgcgcacccc tgagagatgg ttggtgccat gtggaagggtg 120
attgtttcgc tggctcctggt gatgcctggc ccctgtgatg ggctgtttcg ctccctatac 180
agaagtgttt ccatgccacc taaggagagc tcaggacagc cattatttct cacccttac 240
attgaagctg ggaagatcca aaaaggaaga gaattgagtt tggtcggccc tttcccagga 300
ctgaacatga agagttatgc cggcttcctc accgtgaata agacttacaa cagcaacctc 360
ttcttctggt tcttcccagc tcagatacag ccagaagatg cccagtagt tctctggcta 420
cagggtgggc cgggaggttc atccatgttt ggactctttg tggaacatgg gccttatggt 480
gtcacaagta acatgacctt gcgtgacaga gacttcccct ggaccacaac gctctccatg 540
ctttacattg acaatccagt gggcacaggc ttcagtttta ctgatgatac ccacggatat 600
gcagtcaatg aggacgatgt agcacgggat ttatacagtg cactaattca gtttttccag 660
atatttcctg aatataaaaa taatgacttt tatgtcactg gggagtctta tgcagggaaa 720
tatgtgccag ccattgcaca cctcatccat tccctcaacc ctgtgagaga ggtgaagatc 780
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aagcagtgcc atgaatgcat agaacacatc aggaagcaga actggtttga ggcctttgaa 960
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tttaatgatg gaactatagt tgaaaagtac ttgcgagaag atacagtaca gtcagttaag 1200
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caggaatata agaaggcaga aaaaaaagtt tggaagatct ttaaatctga cagtgaagtg 1380
gctggttaca tccggcaagc gggtgacttc catcaggtaa ttattcgagg tggaggacat 1440
attttaccct atgaccagcc tctgagagct tttgacatga ttaatcgatt catttatgga 1500
aaaggatggg atccttatgt tggataaact accttcccaa aagagaacat cagagggttt 1560

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cattgctgaa aagaaaatcg taaaaacaga aaatgtcata ggaataaaaa aattatcttt 1620
tcatatctgc aagatttttt tcatcaataa aaattatcct tgaaacaagt gagcttttgt 1680
ttttgggggg agatgtttac tacaaaatta acatgagtac atgagtaaga attacattat 1740
ttaacttaaa ggatgaaagg tatggatgat gtgacactga gacaagatgt ataaatgaaa 1800
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gtgtttggaa atattattgg ataagaatag ctcaattatc ccaaataaat ggatgaagct 1980
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gaaataaaaa tattatatat aaaagtaaaa aaaaaa 2076

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<210> 164
<211> 476
<212> PRT
<213> Homo sapiens

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<400> 164
Met Val Gly Ala Met Trp Lys Val Ile Val Ser Leu Val Leu Leu Met
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Pro Gly Pro Cys Asp Gly Leu Phe Arg Ser Leu Tyr Arg Ser Val Ser
      20              25              30

Met Pro Pro Lys Gly Asp Ser Gly Gln Pro Leu Phe Leu Thr Pro Tyr
      35              40              45

Ile Glu Ala Gly Lys Ile Gln Lys Gly Arg Glu Leu Ser Leu Val Gly
      50              55              60

Pro Phe Pro Gly Leu Asn Met Lys Ser Tyr Ala Gly Phe Leu Thr Val
      65              70              75              80

Asn Lys Thr Tyr Asn Ser Asn Leu Phe Phe Trp Phe Phe Pro Ala Gln
      85              90              95

Ile Gln Pro Glu Asp Ala Pro Val Val Leu Trp Leu Gln Gly Gly Pro
      100             105             110

Gly Gly Ser Ser Met Phe Gly Leu Phe Val Glu His Gly Pro Tyr Val
      115             120             125

Val Thr Ser Asn Met Thr Leu Arg Asp Arg Asp Phe Pro Trp Thr Thr
      130             135             140

Thr Leu Ser Met Leu Tyr Ile Asp Asn Pro Val Gly Thr Gly Phe Ser
      145             150             155             160

Phe Thr Asp Asp Thr His Gly Tyr Ala Val Asn Glu Asp Asp Val Ala
      165             170             175

Arg Asp Leu Tyr Ser Ala Leu Ile Gln Phe Phe Gln Ile Phe Pro Glu
      180             185             190

Tyr Lys Asn Asn Asp Phe Tyr Val Thr Gly Glu Ser Tyr Ala Gly Lys
      195             200             205

Tyr Val Pro Ala Ile Ala His Leu Ile His Ser Leu Asn Pro Val Arg
      210             215             220

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| | | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Val | Lys | Ile | Asn | Leu | Asn | Gly | Ile | Ala | Ile | Gly | Asp | Gly | Tyr | Ser | 225 | 230 | 235 | 240 |
| Asp | Pro | Glu | Ser | Ile | Ile | Gly | Gly | Tyr | Ala | Glu | Phe | Leu | Tyr | Gln | Ile | 245 | 250 | 255 | |
| Gly | Leu | Leu | Asp | Glu | Lys | Gln | Lys | Lys | Tyr | Phe | Gln | Lys | Gln | Cys | His | 260 | 265 | 270 | |
| Glu | Cys | Ile | Glu | His | Ile | Arg | Lys | Gln | Asn | Trp | Phe | Glu | Ala | Phe | Glu | 275 | 280 | 285 | |
| Ile | Leu | Asp | Lys | Leu | Leu | Asp | Gly | Asp | Leu | Thr | Ser | Asp | Pro | Ser | Tyr | 290 | 295 | 300 | |
| Phe | Gln | Asn | Val | Thr | Gly | Cys | Ser | Asn | Tyr | Tyr | Asn | Phe | Leu | Arg | Cys | 305 | 310 | 315 | 320 |
| Thr | Glu | Pro | Glu | Asp | Gln | Leu | Tyr | Tyr | Val | Lys | Phe | Leu | Ser | Leu | Pro | 325 | 330 | 335 | |
| Glu | Val | Arg | Gln | Ala | Ile | His | Val | Gly | Asn | Gln | Thr | Phe | Asn | Asp | Gly | 340 | 345 | 350 | |
| Thr | Ile | Val | Glu | Lys | Tyr | Leu | Arg | Glu | Asp | Thr | Val | Gln | Ser | Val | Lys | 355 | 360 | 365 | |
| Pro | Trp | Leu | Thr | Glu | Ile | Met | Asn | Asn | Tyr | Lys | Val | Leu | Ile | Tyr | Asn | 370 | 375 | 380 | |
| Gly | Gln | Leu | Asp | Ile | Ile | Val | Ala | Ala | Ala | Leu | Thr | Glu | Arg | Ser | Leu | 385 | 390 | 395 | 400 |
| Met | Gly | Met | Asp | Trp | Lys | Gly | Ser | Gln | Glu | Tyr | Lys | Lys | Ala | Glu | Lys | 405 | 410 | 415 | |
| Lys | Val | Trp | Lys | Ile | Phe | Lys | Ser | Asp | Ser | Glu | Val | Ala | Gly | Tyr | Ile | 420 | 425 | 430 | |
| Arg | Gln | Ala | Gly | Asp | Phe | His | Gln | Val | Ile | Ile | Arg | Gly | Gly | Gly | His | 435 | 440 | 445 | |
| Ile | Leu | Pro | Tyr | Asp | Gln | Pro | Leu | Arg | Ala | Phe | Asp | Met | Ile | Asn | Arg | 450 | 455 | 460 | |
| Phe | Ile | Tyr | Gly | Lys | Gly | Trp | Asp | Pro | Tyr | Val | Gly | | | | | 465 | 470 | 475 | |

<210> 165

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 165
ttccatgccca cctaagggag actc 24

<210> 166
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 166
tggatgaggt gtgcaatggc tggc 24

<210> 167
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 167
agctctcaga ggctgggtcat aggg 24

<210> 168
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 168
gtcggccctt tcccaggact gaacatgaag agttatgccg gcttcctcac 50

<210> 169
<211> 2477
<212> DNA
<213> Homo sapiens

<400> 169
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tcttgctgga gaagaaaggg ctgagggcag agcagggcac tctcactcag ggtgaccagc 180
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gatggaagtc taaaatagga aggaattttg tgtgcaatat cagactctgg gagcagttga 300
cctggagagc ctggggggagg gcctgcctaa caagctttca aaaaacagga gcgacttcca 360
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cttaaggggc cagaaataga gatgctttgt aaaataaaat tttaaaaaaa gcaagtattt 540
tatagcataa aggctagaga ccaaaataga taacaggatt ccctgaacat tcctaagagg 600
gagaaagtat gttaaaaata gaaaaaccaa aatgcagaag gaggagactc acagagctaa 660

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accaggatgg ggaccctggg tcaggccagc ctctttgctc ctcccggaaa ttatTTTTtg 720
tctgaccact ctgccttggt ttttgagaa tcatgtgagg gccaacggg gaaggtggag 780
cagatgagca cacacaggag ccgtctctc accgccggc ctctcagcat ggaacagagg 840
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gtaaggaatg caagcgtatt tcaatatTTc ccaaacttta agaaaaaact ttaagaagg 2460
acatctgcaa aagcaaa 2477

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<210> 170
 <211> 552
 <212> PRT
 <213> Homo sapiens

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<400> 170
Met Gly Thr Leu Gly Gln Ala Ser Leu Phe Ala Pro Pro Gly Asn Tyr
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Phe Trp Ser Asp His Ser Ala Leu Cys Phe Ala Glu Ser Cys Glu Gly
      20              25              30

Gln Pro Gly Lys Val Glu Gln Met Ser Thr His Arg Ser Arg Leu Leu
      35              40              45

Thr Ala Ala Pro Leu Ser Met Glu Gln Arg Gln Pro Trp Pro Arg Ala
      50              55              60

Leu Glu Val Asp Ser Arg Ser Val Val Leu Leu Ser Val Val Trp Val
      65              70              75              80

Leu Leu Ala Pro Pro Ala Ala Gly Met Pro Gln Phe Ser Thr Phe His
      85              90              95

Ser Glu Asn Arg Asp Trp Thr Phe Asn His Leu Thr Val His Gln Gly

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| 100 | | | | | 105 | | | | | 110 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Thr | Gly | Ala | Val | Tyr | Val | Gly | Ala | Ile | Asn | Arg | Val | Tyr | Lys | Leu | Thr |
| | 115 | | | | | | 120 | | | | | 125 | | | |
| Gly | Asn | Leu | Thr | Ile | Gln | Val | Ala | His | Lys | Thr | Gly | Pro | Glu | Glu | Asp |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| Asn | Lys | Ser | Arg | Tyr | Pro | Pro | Leu | Ile | Val | Gln | Pro | Cys | Ser | Glu | Val |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Leu | Thr | Leu | Thr | Asn | Asn | Val | Asn | Lys | Leu | Leu | Ile | Ile | Asp | Tyr | Ser |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Glu | Asn | Arg | Leu | Leu | Ala | Cys | Gly | Ser | Leu | Tyr | Gln | Gly | Val | Cys | Lys |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Leu | Leu | Arg | Leu | Asp | Asp | Leu | Phe | Ile | Leu | Val | Glu | Pro | Ser | His | Lys |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Lys | Glu | His | Tyr | Leu | Ser | Ser | Val | Asn | Lys | Thr | Gly | Thr | Met | Tyr | Gly |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Val | Ile | Val | Arg | Ser | Glu | Gly | Glu | Asp | Gly | Lys | Leu | Phe | Ile | Gly | Thr |
| 225 | | | | | | 230 | | | | | 235 | | | | 240 |
| Ala | Val | Asp | Gly | Lys | Gln | Asp | Tyr | Phe | Pro | Thr | Leu | Ser | Ser | Arg | Lys |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Leu | Pro | Arg | Asp | Pro | Glu | Ser | Ser | Ala | Met | Leu | Asp | Tyr | Glu | Leu | His |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Ser | Asp | Phe | Val | Ser | Ser | Leu | Ile | Lys | Ile | Pro | Ser | Asp | Thr | Leu | Ala |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Leu | Val | Ser | His | Phe | Asp | Ile | Phe | Tyr | Ile | Tyr | Gly | Phe | Ala | Ser | Gly |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Gly | Phe | Val | Tyr | Phe | Leu | Thr | Val | Gln | Pro | Glu | Thr | Pro | Glu | Gly | Val |
| 305 | | | | | | 310 | | | | | 315 | | | | 320 |
| Ala | Ile | Asn | Ser | Ala | Gly | Asp | Leu | Phe | Tyr | Thr | Ser | Arg | Ile | Val | Arg |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Leu | Cys | Lys | Asp | Asp | Pro | Lys | Phe | His | Ser | Tyr | Val | Ser | Leu | Pro | Phe |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| Gly | Cys | Thr | Arg | Ala | Gly | Val | Glu | Tyr | Arg | Leu | Leu | Gln | Ala | Ala | Tyr |
| | | 355 | | | | | 360 | | | | | 365 | | | |
| Leu | Ala | Lys | Pro | Gly | Asp | Ser | Leu | Ala | Gln | Ala | Phe | Asn | Ile | Thr | Ser |
| | 370 | | | | | 375 | | | | | 380 | | | | |
| Gln | Asp | Asp | Val | Leu | Phe | Ala | Ile | Phe | Ser | Lys | Gly | Gln | Lys | Gln | Tyr |
| 385 | | | | | | 390 | | | | | 395 | | | | 400 |
| His | His | Pro | Pro | Asp | Asp | Ser | Ala | Leu | Cys | Ala | Phe | Pro | Ile | Arg | Ala |

| 405 | | | | | 410 | | | | | 415 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Asn | Leu | Gln | Ile | Lys | Glu | Arg | Leu | Gln | Ser | Cys | Tyr | Gln | Gly | Glu |
| | 420 | | | | | | | 425 | | | | | 430 | | |
| Gly | Asn | Leu | Glu | Leu | Asn | Trp | Leu | Leu | Gly | Lys | Asp | Val | Gln | Cys | Thr |
| | 435 | | | | | | 440 | | | | | 445 | | | |
| Lys | Ala | Pro | Val | Pro | Ile | Asp | Asp | Asn | Phe | Cys | Gly | Leu | Asp | Ile | Asn |
| | 450 | | | | | 455 | | | | | 460 | | | | |
| Gln | Pro | Leu | Gly | Gly | Ser | Thr | Pro | Val | Glu | Gly | Leu | Thr | Leu | Tyr | Thr |
| 465 | | | | | | 470 | | | | | 475 | | | | 480 |
| Thr | Ser | Arg | Asp | Arg | Met | Thr | Ser | Val | Ala | Ser | Tyr | Val | Tyr | Asn | Gly |
| | | | | 485 | | | | | 490 | | | | | 495 | |
| Tyr | Ser | Val | Val | Phe | Val | Gly | Thr | Lys | Ser | Gly | Lys | Leu | Lys | Lys | Val |
| | | | 500 | | | | | 505 | | | | | 510 | | |
| Arg | Val | Tyr | Glu | Phe | Arg | Cys | Ser | Asn | Ala | Ile | His | Leu | Leu | Ser | Lys |
| | 515 | | | | | | 520 | | | | | 525 | | | |
| Glu | Ser | Leu | Leu | Glu | Gly | Ser | Tyr | Trp | Trp | Arg | Phe | Asn | Tyr | Arg | Gln |
| | 530 | | | | | 535 | | | | | 540 | | | | |
| Leu | Tyr | Phe | Leu | Gly | Glu | Gln | Arg | | | | | | | | |
| 545 | | | | | 550 | | | | | | | | | | |

<210> 171
 <211> 20
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 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 171
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20

<210> 172
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 172
 cttctgccct ttggagaaga tggc

24

<210> 173
 <211> 43
 <212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 173

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43

<210> 174

<211> 3106

<212> DNA

<213> Homo sapiens

<220>

<221> modified_base

<222> (1683)..(1683)

<223> a, t, c or g

<400> 174

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| aacacgcgat | gaccacgtgg | agcctccggc | ggaggccggc | ccgcacgctg | ggactcctgc | 120 |
| tgctgggtcgt | cttgggcttc | ctgggtgctcc | gcaggctgga | ctggagcacc | ctggtccttc | 180 |
| tgcggctccg | ccatcgacag | ctggggctgc | aggccaagg | ctggaacttc | atgctggagg | 240 |
| attccacctt | ctggatcttc | gggggtcca | tccactattt | ccgtgtgccc | aggagtagt | 300 |
| ggagggaccg | cctgctgaag | atgaaggcct | gtggcttgaa | caccctcacc | acctatgttc | 360 |
| cgtggaacct | gcatgagcca | gaaagaggca | aatttgactt | ctctgggaac | ctggacctgg | 420 |
| aggccttcgt | cctgatggcc | gcagagatcg | ggctgtgggt | gattctgcgt | ccaggccccct | 480 |
| acatctgcag | tgagatggac | ctcgggggct | tgcccagctg | gctactccaa | gaccttgcca | 540 |
| tgaggctgag | gacaacttac | aagggttca | ccgaagcagt | ggacctttat | tttgaccacc | 600 |
| tgatgtccag | ggtggtgcc | ctccagtaca | agcgtggggg | acctatcatt | gccgtgcagg | 660 |
| tggagaatga | atatggttcc | tataataaag | acccgcata | catgccctac | gtcaagaagg | 720 |
| cactggagga | ccgtggcatt | gtggaactgc | tcttgacttc | agacaacaag | gatgggctga | 780 |
| gcaaggggat | tgtccaggga | gtcttgcca | ccatcaactt | gcagtcaaca | cacgagctgc | 840 |
| agctactgac | cacctttctc | ttcaacgtcc | aggggactca | gccaagatg | gtgatggagt | 900 |
| actggacggg | gtggtttgac | tctgtgggag | gccctcaca | tatcttggat | tcttctgagg | 960 |
| ttttgaaaac | cgtgtctgcc | attgtggacg | ccggctcctc | catcaacctc | tacatgttcc | 1020 |
| acggaggcac | caactttggc | ttcatgaatg | gagccatgca | cttccatgac | tacaagtcag | 1080 |
| atgtcaccag | ctatgactat | gatgctgtgc | tgacagaagc | cggcgattac | acggccaagt | 1140 |
| acatgaagct | tcgagacttc | ttcggtcca | tctcaggcat | ccctctccct | ccccacctg | 1200 |
| accttcttcc | caagatgccg | tatgagccct | taacgccagt | cttgtacctg | tctctgtggg | 1260 |
| acgccctcaa | gtacctgggg | gagccaatca | agtctgaaaa | gcccataaac | atggagaacc | 1320 |
| tgccagtcaa | tgggggaaat | ggacagtcct | tcgggtacat | tctctatgag | accagcatca | 1380 |
| cctcgtctgg | catcctcagt | ggccacgtgc | atgatcgggg | gcagggtgtt | gtgaacacag | 1440 |
| tatccatagg | attcttggac | tacaagacaa | cgaagattgc | tgtccccctg | atccagggtt | 1500 |
| acaccgtgct | gaggatcttg | gtggagaatc | gtgggcgagt | caactatggg | gagaatattg | 1560 |
| atgaccagcg | caaaggctta | attggaaatc | tctatctgaa | tgattcacc | ctgaaaaact | 1620 |
| tcagaatcta | tagcctggat | atgaagaaga | gcttctttca | gaggttcggc | ctggacaaat | 1680 |
| gngttccct | cccagaaaca | cccacattac | ctgctttctt | cttgggtagc | ttgtccatca | 1740 |
| gtccacgcgc | ttgtgacacc | tttctgaagc | tggagggtcg | ggagaagggg | gttgtattca | 1800 |
| tcaatggcca | gaaccttggg | cgttactgga | acattggacc | ccagaagacg | ctttacctcc | 1860 |
| caggtcctcg | gttgagcagc | ggaatcaacc | aggtcatcgt | ttttgaggag | acgatggcgg | 1920 |
| gccctgcatt | acagttcacg | gaaaccccc | acctgggcag | gaaccagtac | attaagttag | 1980 |
| cgggtggcacc | ccctcctgct | ggtgccagtg | ggagactgcc | gcctcctctt | gacctgaagc | 2040 |
| ctggtggctg | ctgccccacc | cctcactgca | aaagcatctc | cttaagtagc | aacctcaggg | 2100 |
| actgggggct | acagtctgcc | cctgtctcag | ctcaaaacct | taagcctgca | gggaaagggtg | 2160 |
| ggatggctct | gggcctggct | ttgttgatga | tggctttcct | acagccctgc | tcttgtgccg | 2220 |

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aggtgtcgg gctgtctcta ggggtgggagc agctaatacag atcgcccagc ctttggccct 2280
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gactcaggcg tgctctttgc tggttcctgg gaggttggc cacatccctc atggcccat 2400
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<210> 175
 <211> 636
 <212> PRT
 <213> Homo sapiens

<220>
 <221> MOD_RES
 <222> (539)
 <223> Any amino acid

<400> 175

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Thr | Thr | Trp | Ser | Leu | Arg | Arg | Arg | Pro | Ala | Arg | Thr | Leu | Gly | Leu |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Leu | Leu | Leu | Val | Val | Leu | Gly | Phe | Leu | Val | Leu | Arg | Arg | Leu | Asp | Trp |
| | | | 20 | | | | | 25 | | | | | | 30 | |
| Ser | Thr | Leu | Val | Pro | Leu | Arg | Leu | Arg | His | Arg | Gln | Leu | Gly | Leu | Gln |
| | | | 35 | | | | | 40 | | | | | 45 | | |
| Ala | Lys | Gly | Trp | Asn | Phe | Met | Leu | Glu | Asp | Ser | Thr | Phe | Trp | Ile | Phe |
| | | | 50 | | | | 55 | | | | | 60 | | | |
| Gly | Gly | Ser | Ile | His | Tyr | Phe | Arg | Val | Pro | Arg | Glu | Tyr | Trp | Arg | Asp |
| | | | 65 | | | 70 | | | | 75 | | | | | 80 |
| Arg | Leu | Leu | Lys | Met | Lys | Ala | Cys | Gly | Leu | Asn | Thr | Leu | Thr | Thr | Tyr |
| | | | | 85 | | | | | 90 | | | | | | 95 |
| Val | Pro | Trp | Asn | Leu | His | Glu | Pro | Glu | Arg | Gly | Lys | Phe | Asp | Phe | Ser |
| | | | 100 | | | | | | 105 | | | | | 110 | |
| Gly | Asn | Leu | Asp | Leu | Glu | Ala | Phe | Val | Leu | Met | Ala | Ala | Glu | Ile | Gly |
| | | | 115 | | | | 120 | | | | | 125 | | | |
| Leu | Trp | Val | Ile | Leu | Arg | Pro | Gly | Pro | Tyr | Ile | Cys | Ser | Glu | Met | Asp |
| | | | 130 | | | | 135 | | | | | 140 | | | |
| Leu | Gly | Gly | Leu | Pro | Ser | Trp | Leu | Leu | Gln | Asp | Pro | Gly | Met | Arg | Leu |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Thr | Thr | Tyr | Lys | Gly | Phe | Thr | Glu | Ala | Val | Asp | Leu | Tyr | Phe | Asp | 165 | 170 | 175 |
| His | Leu | Met | Ser | Arg | Val | Val | Pro | Leu | Gln | Tyr | Lys | Arg | Gly | Gly | Pro | 180 | 185 | 190 |
| Ile | Ile | Ala | Val | Gln | Val | Glu | Asn | Glu | Tyr | Gly | Ser | Tyr | Asn | Lys | Asp | 195 | 200 | 205 |
| Pro | Ala | Tyr | Met | Pro | Tyr | Val | Lys | Lys | Ala | Leu | Glu | Asp | Arg | Gly | Ile | 210 | 215 | 220 |
| Val | Glu | Leu | Leu | Leu | Thr | Ser | Asp | Asn | Lys | Asp | Gly | Leu | Ser | Lys | Gly | 225 | 230 | 235 |
| Ile | Val | Gln | Gly | Val | Leu | Ala | Thr | Ile | Asn | Leu | Gln | Ser | Thr | His | Glu | 245 | 250 | 255 |
| Leu | Gln | Leu | Leu | Thr | Thr | Phe | Leu | Phe | Asn | Val | Gln | Gly | Thr | Gln | Pro | 260 | 265 | 270 |
| Lys | Met | Val | Met | Glu | Tyr | Trp | Thr | Gly | Trp | Phe | Asp | Ser | Trp | Gly | Gly | 275 | 280 | 285 |
| Pro | His | Asn | Ile | Leu | Asp | Ser | Ser | Glu | Val | Leu | Lys | Thr | Val | Ser | Ala | 290 | 295 | 300 |
| Ile | Val | Asp | Ala | Gly | Ser | Ser | Ile | Asn | Leu | Tyr | Met | Phe | His | Gly | Gly | 305 | 310 | 315 |
| Thr | Asn | Phe | Gly | Phe | Met | Asn | Gly | Ala | Met | His | Phe | His | Asp | Tyr | Lys | 325 | 330 | 335 |
| Ser | Asp | Val | Thr | Ser | Tyr | Asp | Tyr | Asp | Ala | Val | Leu | Thr | Glu | Ala | Gly | 340 | 345 | 350 |
| Asp | Tyr | Thr | Ala | Lys | Tyr | Met | Lys | Leu | Arg | Asp | Phe | Phe | Gly | Ser | Ile | 355 | 360 | 365 |
| Ser | Gly | Ile | Pro | Leu | Pro | Pro | Pro | Pro | Asp | Leu | Leu | Pro | Lys | Met | Pro | 370 | 375 | 380 |
| Tyr | Glu | Pro | Leu | Thr | Pro | Val | Leu | Tyr | Leu | Ser | Leu | Trp | Asp | Ala | Leu | 385 | 390 | 395 |
| Lys | Tyr | Leu | Gly | Glu | Pro | Ile | Lys | Ser | Glu | Lys | Pro | Ile | Asn | Met | Glu | 405 | 410 | 415 |
| Asn | Leu | Pro | Val | Asn | Gly | Gly | Asn | Gly | Gln | Ser | Phe | Gly | Tyr | Ile | Leu | 420 | 425 | 430 |
| Tyr | Glu | Thr | Ser | Ile | Thr | Ser | Ser | Gly | Ile | Leu | Ser | Gly | His | Val | His | 435 | 440 | 445 |
| Asp | Arg | Gly | Gln | Val | Phe | Val | Asn | Thr | Val | Ser | Ile | Gly | Phe | Leu | Asp | 450 | 455 | 460 |

Tyr Lys Thr Thr Lys Ile Ala Val Pro Leu Ile Gln Gly Tyr Thr Val
 465 470 475 480
 Leu Arg Ile Leu Val Glu Asn Arg Gly Arg Val Asn Tyr Gly Glu Asn
 485 490 495
 Ile Asp Asp Gln Arg Lys Gly Leu Ile Gly Asn Leu Tyr Leu Asn Asp
 500 505 510
 Ser Pro Leu Lys Asn Phe Arg Ile Tyr Ser Leu Asp Met Lys Lys Ser
 515 520 525
 Phe Phe Gln Arg Phe Gly Leu Asp Lys Trp Xaa Ser Leu Pro Glu Thr
 530 535 540
 Pro Thr Leu Pro Ala Phe Phe Leu Gly Ser Leu Ser Ile Ser Ser Thr
 545 550 555 560
 Pro Cys Asp Thr Phe Leu Lys Leu Glu Gly Trp Glu Lys Gly Val Val
 565 570 575
 Phe Ile Asn Gly Gln Asn Leu Gly Arg Tyr Trp Asn Ile Gly Pro Gln
 580 585 590
 Lys Thr Leu Tyr Leu Pro Gly Pro Trp Leu Ser Ser Gly Ile Asn Gln
 595 600 605
 Val Ile Val Phe Glu Glu Thr Met Ala Gly Pro Ala Leu Gln Phe Thr
 610 615 620
 Glu Thr Pro His Leu Gly Arg Asn Gln Tyr Ile Lys
 625 630 635

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 <211> 2505
 <212> DNA
 <213> Homo sapiens

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 aaggggagca aagccgggct cggcccagg cccccaggac ctccatctcc caatgttgga 180
 ggaatccgac acgtgacggt ctgtccgccc tctcagacta gaggagcgct gtaaaccgcca 240
 tggtcccaa gaagctgtcc tgcttcggtt ccctgctgct gccgctcagc ctgacgctac 300
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 atgtgccctg gaactaccac gagccacagc ctgggggtcta taactttaat ggcagccggg 540
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 gaccttacat ctgtgcagag tgggagatgg ggggtctccc atcctggttg cttcgaaaac 660
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 ctgacaacat gaccaaaatc tttaccctgc ttcggaagta tgaaccccat gggccattgg 1020
 taaactctga gtactacaca ggctggctgg attactgggg ccagaatcac tccacacggt 1080

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ctgtgtcagc tgaacacaaa ggactagaga acatgctcaa gttgggagcc agtgtgaaca 1140
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<210> 177
 <211> 654
 <212> PRT
 <213> Homo sapiens

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<400> 177
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Ser Leu Thr Leu Leu Leu Pro Gln Ala Asp Thr Arg Ser Phe Val Val
      20              25              30

Asp Arg Gly His Asp Arg Phe Leu Leu Asp Gly Ala Pro Phe Arg Tyr
      35              40              45

Val Ser Gly Ser Leu His Tyr Phe Arg Val Pro Arg Val Leu Trp Ala
      50              55              60

Asp Arg Leu Leu Lys Met Arg Trp Ser Gly Leu Asn Ala Ile Gln Phe
      65              70              75              80

Tyr Val Pro Trp Asn Tyr His Glu Pro Gln Pro Gly Val Tyr Asn Phe
      85              90              95

Asn Gly Ser Arg Asp Leu Ile Ala Phe Leu Asn Glu Ala Ala Leu Ala
      100             105             110

Asn Leu Leu Val Ile Leu Arg Pro Gly Pro Tyr Ile Cys Ala Glu Trp
      115             120             125

Glu Met Gly Gly Leu Pro Ser Trp Leu Leu Arg Lys Pro Glu Ile His
      130             135             140

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| | | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Arg | Thr | Ser | Asp | Pro | Asp | Phe | Leu | Ala | Ala | Val | Asp | Ser | Trp | Phe | 145 | 150 | 155 | 160 |
| Lys | Val | Leu | Leu | Pro | Lys | Ile | Tyr | Pro | Trp | Leu | Tyr | His | Asn | Gly | Gly | 165 | 170 | 175 | |
| Asn | Ile | Ile | Ser | Ile | Gln | Val | Glu | Asn | Glu | Tyr | Gly | Ser | Tyr | Arg | Ala | 180 | 185 | 190 | |
| Cys | Asp | Phe | Ser | Tyr | Met | Arg | His | Leu | Ala | Gly | Leu | Phe | Arg | Ala | Leu | 195 | 200 | 205 | |
| Leu | Gly | Glu | Lys | Ile | Leu | Leu | Phe | Thr | Thr | Asp | Gly | Pro | Glu | Gly | Leu | 210 | 215 | 220 | |
| Lys | Cys | Gly | Ser | Leu | Arg | Gly | Leu | Tyr | Thr | Thr | Val | Asp | Phe | Gly | Pro | 225 | 230 | 235 | 240 |
| Ala | Asp | Asn | Met | Thr | Lys | Ile | Phe | Thr | Leu | Leu | Arg | Lys | Tyr | Glu | Pro | 245 | 250 | 255 | |
| His | Gly | Pro | Leu | Val | Asn | Ser | Glu | Tyr | Tyr | Thr | Gly | Trp | Leu | Asp | Tyr | 260 | 265 | 270 | |
| Trp | Gly | Gln | Asn | His | Ser | Thr | Arg | Ser | Val | Ser | Ala | Val | Thr | Lys | Gly | 275 | 280 | 285 | |
| Leu | Glu | Asn | Met | Leu | Lys | Leu | Gly | Ala | Ser | Val | Asn | Met | Tyr | Met | Phe | 290 | 295 | 300 | |
| His | Gly | Gly | Thr | Asn | Phe | Gly | Tyr | Trp | Asn | Gly | Ala | Asp | Lys | Lys | Gly | 305 | 310 | 315 | 320 |
| Arg | Phe | Leu | Pro | Ile | Thr | Thr | Ser | Tyr | Asp | Tyr | Asp | Ala | Pro | Ile | Ser | 325 | 330 | 335 | |
| Glu | Ala | Gly | Asp | Pro | Thr | Pro | Lys | Leu | Phe | Ala | Leu | Arg | Asp | Val | Ile | 340 | 345 | 350 | |
| Ser | Lys | Phe | Gln | Glu | Val | Pro | Leu | Gly | Pro | Leu | Pro | Pro | Pro | Ser | Pro | 355 | 360 | 365 | |
| Lys | Met | Met | Leu | Gly | Pro | Val | Thr | Leu | His | Leu | Val | Gly | His | Leu | Leu | 370 | 375 | 380 | |
| Ala | Phe | Leu | Asp | Leu | Leu | Cys | Pro | Arg | Gly | Pro | Ile | His | Ser | Ile | Leu | 385 | 390 | 395 | 400 |
| Pro | Met | Thr | Phe | Glu | Ala | Val | Lys | Gln | Asp | His | Gly | Phe | Met | Leu | Tyr | 405 | 410 | 415 | |
| Arg | Thr | Tyr | Met | Thr | His | Thr | Ile | Phe | Glu | Pro | Thr | Pro | Phe | Trp | Val | 420 | 425 | 430 | |
| Pro | Asn | Asn | Gly | Val | His | Asp | Arg | Ala | Tyr | Val | Met | Val | Asp | Gly | Val | 435 | 440 | 445 | |

Phe Gln Gly Val Val Glu Arg Asn Met Arg Asp Lys Leu Phe Leu Thr
450 455 460
Gly Lys Leu Gly Ser Lys Leu Asp Ile Leu Val Glu Asn Met Gly Arg
465 470 475 480
Leu Ser Phe Gly Ser Asn Ser Ser Asp Phe Lys Gly Leu Leu Lys Pro
485 490 495
Pro Ile Leu Gly Gln Thr Ile Leu Thr Gln Trp Met Met Phe Pro Leu
500 505 510
Lys Ile Asp Asn Leu Val Lys Trp Trp Phe Pro Leu Gln Leu Pro Lys
515 520 525
Trp Pro Tyr Pro Gln Ala Pro Ser Gly Pro Thr Phe Tyr Ser Lys Thr
530 535 540
Phe Pro Ile Leu Gly Ser Val Gly Asp Thr Phe Leu Tyr Leu Pro Gly
545 550 555 560
Trp Thr Lys Gly Gln Val Trp Ile Asn Gly Phe Asn Leu Gly Arg Tyr
565 570 575
Trp Thr Lys Gln Gly Pro Gln Gln Thr Leu Tyr Val Pro Arg Phe Leu
580 585 590
Leu Phe Pro Arg Gly Ala Leu Asn Lys Ile Thr Leu Leu Glu Leu Glu
595 600 605
Asp Val Pro Leu Gln Pro Gln Val Gln Phe Leu Asp Lys Pro Ile Leu
610 615 620
Asn Ser Thr Ser Thr Leu His Arg Thr His Ile Asn Ser Leu Ser Ala
625 630 635 640
Asp Thr Leu Ser Ala Ser Glu Pro Met Glu Leu Ser Gly His
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<210> 178

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 178

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24

<210> 179

<211> 24

<212> DNA

<213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 179
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<210> 180
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 180
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<210> 181
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 181
 ccagctatga ctatgatgca cc 22

<210> 182
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 182
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<210> 183
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 183
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<210> 184
 <211> 1947

<212> DNA
<213> Homo sapiens

<400> 184

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<210> 185
<211> 501
<212> PRT
<213> Homo sapiens

<400> 185

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  20              25             30

Ile Pro Leu Lys Glu Tyr Ser Phe Glu Lys Val Arg Glu Glu Ser Ser
  35              40             45

Phe Ser Asp Ile Pro Asp Val Lys Asn Asp Phe Ala Phe Leu Leu His
  50              55             60

Met Val Asp Gln Tyr Asp Gln Leu Tyr Ser Lys Arg Phe Gly Val Phe
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| | | | | | | | | | | | | | | | |
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| Leu | Ser | Glu | Val | Ser | Glu | Asn | Lys | Leu | Arg | Glu | Ile | Ser | Leu | Asn | His |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| Glu | Trp | Thr | Phe | Glu | Lys | Leu | Arg | Gln | His | Ile | Ser | Arg | Asn | Ala | Gln |
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| Asp | Lys | Gln | Glu | Leu | His | Leu | Phe | Met | Leu | Ser | Gly | Val | Pro | Asp | Ala |
| | | 115 | | | | | 120 | | | | | 125 | | | |
| Val | Phe | Asp | Leu | Thr | Asp | Leu | Asp | Val | Leu | Lys | Leu | Glu | Leu | Ile | Pro |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| Glu | Ala | Lys | Ile | Pro | Ala | Lys | Ile | Ser | Gln | Met | Thr | Asn | Leu | Gln | Glu |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Leu | His | Leu | Cys | His | Cys | Pro | Ala | Lys | Val | Glu | Gln | Thr | Ala | Phe | Ser |
| | | | 165 | | | | | | 170 | | | | | 175 | |
| Phe | Leu | Arg | Asp | His | Leu | Arg | Cys | Leu | His | Val | Lys | Phe | Thr | Asp | Val |
| | | 180 | | | | | | 185 | | | | | 190 | | |
| Ala | Glu | Ile | Pro | Ala | Trp | Val | Tyr | Leu | Leu | Lys | Asn | Leu | Arg | Glu | Leu |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Tyr | Leu | Ile | Gly | Asn | Leu | Asn | Ser | Glu | Asn | Asn | Lys | Met | Ile | Gly | Leu |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Glu | Ser | Leu | Arg | Glu | Leu | Arg | His | Leu | Lys | Ile | Leu | His | Val | Lys | Ser |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Asn | Leu | Thr | Lys | Val | Pro | Ser | Asn | Ile | Thr | Asp | Val | Ala | Pro | His | Leu |
| | | | 245 | | | | | | 250 | | | | | 255 | |
| Thr | Lys | Leu | Val | Ile | His | Asn | Asp | Gly | Thr | Lys | Leu | Leu | Val | Leu | Asn |
| | | 260 | | | | | | 265 | | | | | 270 | | |
| Ser | Leu | Lys | Lys | Met | Met | Asn | Val | Ala | Glu | Leu | Glu | Leu | Gln | Asn | Cys |
| | 275 | | | | | 280 | | | | | | 285 | | | |
| Glu | Leu | Glu | Arg | Ile | Pro | His | Ala | Ile | Phe | Ser | Leu | Ser | Asn | Leu | Gln |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Glu | Leu | Asp | Leu | Lys | Ser | Asn | Asn | Ile | Arg | Thr | Ile | Glu | Glu | Ile | Ile |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Ser | Phe | Gln | His | Leu | Lys | Arg | Leu | Thr | Cys | Leu | Lys | Leu | Trp | His | Asn |
| | | | 325 | | | | | | 330 | | | | | 335 | |
| Lys | Ile | Val | Thr | Ile | Pro | Pro | Ser | Ile | Thr | His | Val | Lys | Asn | Leu | Glu |
| | | 340 | | | | | | 345 | | | | | 350 | | |
| Ser | Leu | Tyr | Phe | Ser | Asn | Asn | Lys | Leu | Glu | Ser | Leu | Pro | Val | Ala | Val |
| | 355 | | | | | | 360 | | | | | 365 | | | |
| Phe | Ser | Leu | Gln | Lys | Leu | Arg | Cys | Leu | Asp | Val | Ser | Tyr | Asn | Asn | Ile |

| 370 | 375 | 380 |
|---|-----|-------------|
| Ser Met Ile Pro Ile Glu Ile Gly Leu Leu Gln Asn Leu Gln His Leu | | |
| 385 | 390 | 395 400 |
| His Ile Thr Gly Asn Lys Val Asp Ile Leu Pro Lys Gln Leu Phe Lys | | |
| | 405 | 410 415 |
| Cys Ile Lys Leu Arg Thr Leu Asn Leu Gly Gln Asn Cys Ile Thr Ser | | |
| | 420 | 425 430 |
| Leu Pro Glu Lys Val Gly Gln Leu Ser Gln Leu Thr Gln Leu Glu Leu | | |
| | 435 | 440 445 |
| Lys Gly Asn Cys Leu Asp Arg Leu Pro Ala Gln Leu Gly Gln Cys Arg | | |
| | 450 | 455 460 |
| Met Leu Lys Lys Ser Gly Leu Val Val Glu Asp His Leu Phe Asp Thr | | |
| | 465 | 470 475 480 |
| Leu Pro Leu Glu Val Lys Glu Ala Leu Asn Gln Asp Ile Asn Ile Pro | | |
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| Phe Ala Asn Gly Ile | | |
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<210> 186

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 186

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21

<210> 187

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 187

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24

<210> 188

<211> 47

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

oligonucleotide probe

<400> 188

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47

<210> 189

<211> 2917

<212> DNA

<213> Homo sapiens

<400> 189

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<210> 190

<211> 607

<212> PRT

<213> Homo sapiens

<400> 190

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Thr Val Ser Leu Gly Gly Ala Asn Met Ala Glu Thr His Lys Ala Met
35 40 45

Ile Leu Gln Leu Asn Pro Ser Glu Asn Cys Thr Trp Thr Ile Glu Arg
50 55 60

Pro Glu Asn Lys Ser Ile Arg Ile Ile Phe Ser Tyr Val Gln Leu Asp
65 70 75 80

Pro Asp Gly Ser Cys Glu Ser Glu Asn Ile Lys Val Phe Asp Gly Thr
85 90 95

Ser Ser Asn Gly Pro Leu Leu Gly Gln Val Cys Ser Lys Asn Asp Tyr
100 105 110

Val Pro Val Phe Glu Ser Ser Ser Ser Thr Leu Thr Phe Gln Ile Val
115 120 125

Thr Asp Ser Ala Arg Ile Gln Arg Thr Val Phe Val Phe Tyr Tyr Phe
130 135 140

Phe Ser Pro Asn Ile Ser Ile Pro Asn Cys Gly Gly Tyr Leu Asp Thr
145 150 155 160

Leu Glu Gly Ser Phe Thr Ser Pro Asn Tyr Pro Lys Pro His Pro Glu
165 170 175

Leu Ala Tyr Cys Val Trp His Ile Gln Val Glu Lys Asp Tyr Lys Ile
180 185 190

Lys Leu Asn Phe Lys Glu Ile Phe Leu Glu Ile Asp Lys Gln Cys Lys
195 200 205

Phe Asp Phe Leu Ala Ile Tyr Asp Gly Pro Ser Thr Asn Ser Gly Leu
210 215 220

Ile Gly Gln Val Cys Gly Arg Val Thr Pro Thr Phe Glu Ser Ser Ser
225 230 235 240

Asn Ser Leu Thr Val Val Leu Ser Thr Asp Tyr Ala Asn Ser Tyr Arg
245 250 255

| | | | | | | | | | | | | | | | | | | | |
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| Gly | Phe | Ser | Ala | Ser | Tyr | Thr | Ser | Ile | Tyr | Ala | Glu | Asn | Ile | Asn | Thr | 260 | 265 | 270 | |
| Thr | Ser | Leu | Thr | Cys | Ser | Ser | Asp | Arg | Met | Arg | Val | Ile | Ile | Ser | Lys | 275 | 280 | 285 | |
| Ser | Tyr | Leu | Glu | Ala | Phe | Asn | Ser | Asn | Gly | Asn | Asn | Leu | Gln | Leu | Lys | 290 | 295 | 300 | |
| Asp | Pro | Thr | Cys | Arg | Pro | Lys | Leu | Ser | Asn | Val | Val | Glu | Phe | Ser | Val | 305 | 310 | 315 | 320 |
| Pro | Leu | Asn | Gly | Cys | Gly | Thr | Ile | Arg | Lys | Val | Glu | Asp | Gln | Ser | Ile | 325 | 330 | 335 | |
| Thr | Tyr | Thr | Asn | Ile | Ile | Thr | Phe | Ser | Ala | Ser | Ser | Thr | Ser | Glu | Val | 340 | 345 | 350 | |
| Ile | Thr | Arg | Gln | Lys | Gln | Leu | Gln | Ile | Ile | Val | Lys | Cys | Glu | Met | Gly | 355 | 360 | 365 | |
| His | Asn | Ser | Thr | Val | Glu | Ile | Ile | Tyr | Ile | Thr | Glu | Asp | Asp | Val | Ile | 370 | 375 | 380 | |
| Gln | Ser | Gln | Asn | Ala | Leu | Gly | Lys | Tyr | Asn | Thr | Ser | Met | Ala | Leu | Phe | 385 | 390 | 395 | 400 |
| Glu | Ser | Asn | Ser | Phe | Glu | Lys | Thr | Ile | Leu | Glu | Ser | Pro | Tyr | Tyr | Val | 405 | 410 | 415 | |
| Asp | Leu | Asn | Gln | Thr | Leu | Phe | Val | Gln | Val | Ser | Leu | His | Thr | Ser | Asp | 420 | 425 | 430 | |
| Pro | Asn | Leu | Val | Val | Phe | Leu | Asp | Thr | Cys | Arg | Ala | Ser | Pro | Thr | Ser | 435 | 440 | 445 | |
| Asp | Phe | Ala | Ser | Pro | Thr | Tyr | Asp | Leu | Ile | Lys | Ser | Gly | Cys | Ser | Arg | 450 | 455 | 460 | |
| Asp | Glu | Thr | Cys | Lys | Val | Tyr | Pro | Leu | Phe | Gly | His | Tyr | Gly | Arg | Phe | 465 | 470 | 475 | 480 |
| Gln | Phe | Asn | Ala | Phe | Lys | Phe | Leu | Arg | Ser | Met | Ser | Ser | Val | Tyr | Leu | 485 | 490 | 495 | |
| Gln | Cys | Lys | Val | Leu | Ile | Cys | Asp | Ser | Ser | Asp | His | Gln | Ser | Arg | Cys | 500 | 505 | 510 | |
| Asn | Gln | Gly | Cys | Val | Ser | Arg | Ser | Lys | Arg | Asp | Ile | Ser | Ser | Tyr | Lys | 515 | 520 | 525 | |
| Trp | Lys | Thr | Asp | Ser | Ile | Ile | Gly | Pro | Ile | Arg | Leu | Lys | Arg | Asp | Arg | 530 | 535 | 540 | |
| Ser | Ala | Ser | Gly | Asn | Ser | Gly | Phe | Gln | His | Glu | Thr | His | Ala | Glu | Glu | 545 | 550 | 555 | 560 |

Thr Pro Asn Gln Pro Phe Asn Ser Val His Leu Phe Ser Phe Met Val
565 570 575

Leu Ala Leu Asn Val Val Thr Val Ala Thr Ile Thr Val Arg His Phe
580 585 590

Val Asn Gln Arg Ala Asp Tyr Lys Tyr Gln Lys Leu Gln Asn Tyr
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<210> 191

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 191

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21

<210> 192

<211> 22

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

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22

<210> 193

<211> 47

<212> DNA

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<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 193

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47

<210> 194

<211> 2362

<212> DNA

<213> Homo sapiens

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taaactcatt gtgcaaatgt aa
2362

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<210> 195

<211> 467

<212> PRT

<213> Homo sapiens

<400> 195

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1 5 10 15

Leu Leu Leu Leu Leu Pro Pro Pro Pro Cys Pro Ala His Ser Ala Thr
20 25 30

Arg Phe Asp Pro Thr Trp Glu Ser Leu Asp Ala Arg Gln Leu Pro Ala
35 40 45

Trp Phe Asp Gln Ala Lys Phe Gly Ile Phe Ile His Trp Gly Val Phe
50 55 60

Ser Val Pro Ser Phe Gly Ser Glu Trp Phe Trp Trp Tyr Trp Gln Lys
65 70 75 80

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Glu | Lys | Ile | Pro | Lys | Tyr | Val | Glu | Phe | Met | Lys | Asp | Asn | Tyr | Pro | Pro | 85 | 90 | 95 |
| Ser | Phe | Lys | Tyr | Glu | Asp | Phe | Gly | Pro | Leu | Phe | Thr | Ala | Lys | Phe | Phe | 100 | 105 | 110 |
| Asn | Ala | Asn | Gln | Trp | Ala | Asp | Ile | Phe | Gln | Ala | Ser | Gly | Ala | Lys | Tyr | 115 | 120 | 125 |
| Ile | Val | Leu | Thr | Ser | Lys | His | His | Glu | Gly | Phe | Thr | Leu | Trp | Gly | Ser | 130 | 135 | 140 |
| Glu | Tyr | Ser | Trp | Asn | Trp | Asn | Ala | Ile | Asp | Glu | Gly | Pro | Lys | Arg | Asp | 145 | 150 | 155 |
| Ile | Val | Lys | Glu | Leu | Glu | Val | Ala | Ile | Arg | Asn | Arg | Thr | Asp | Leu | Arg | 165 | 170 | 175 |
| Phe | Gly | Leu | Tyr | Tyr | Ser | Leu | Phe | Glu | Trp | Phe | His | Pro | Leu | Phe | Leu | 180 | 185 | 190 |
| Glu | Asp | Glu | Ser | Ser | Ser | Phe | His | Lys | Arg | Gln | Phe | Pro | Val | Ser | Lys | 195 | 200 | 205 |
| Thr | Leu | Pro | Glu | Leu | Tyr | Glu | Leu | Val | Asn | Asn | Tyr | Gln | Pro | Glu | Val | 210 | 215 | 220 |
| Leu | Trp | Ser | Asp | Gly | Asp | Gly | Gly | Ala | Pro | Asp | Gln | Tyr | Trp | Asn | Ser | 225 | 230 | 235 |
| Thr | Gly | Phe | Leu | Ala | Trp | Leu | Tyr | Asn | Glu | Ser | Pro | Val | Arg | Gly | Thr | 245 | 250 | 255 |
| Val | Val | Thr | Asn | Asp | Arg | Trp | Gly | Ala | Gly | Ser | Ile | Cys | Lys | His | Gly | 260 | 265 | 270 |
| Gly | Phe | Tyr | Thr | Cys | Ser | Asp | Arg | Tyr | Asn | Pro | Gly | His | Leu | Leu | Pro | 275 | 280 | 285 |
| His | Lys | Trp | Glu | Asn | Cys | Met | Thr | Ile | Asp | Lys | Leu | Ser | Trp | Gly | Tyr | 290 | 295 | 300 |
| Arg | Arg | Glu | Ala | Gly | Ile | Ser | Asp | Tyr | Leu | Thr | Ile | Glu | Glu | Leu | Val | 305 | 310 | 315 |
| Lys | Gln | Leu | Val | Glu | Thr | Val | Ser | Cys | Gly | Gly | Asn | Leu | Leu | Met | Asn | 325 | 330 | 335 |
| Ile | Gly | Pro | Thr | Leu | Asp | Gly | Thr | Ile | Ser | Val | Val | Phe | Glu | Glu | Arg | 340 | 345 | 350 |
| Leu | Arg | Gln | Val | Gly | Ser | Trp | Leu | Lys | Val | Asn | Gly | Glu | Ala | Ile | Tyr | 355 | 360 | 365 |
| Glu | Thr | Tyr | Thr | Trp | Arg | Ser | Gln | Asn | Asp | Thr | Val | Thr | Pro | Asp | Val | 370 | 375 | 380 |

Trp Tyr Thr Ser Lys Pro Lys Glu Lys Leu Val Tyr Ala Ile Phe Leu
 385 390 395 400
 Lys Trp Pro Thr Ser Gly Gln Leu Phe Leu Gly His Pro Lys Ala Ile
 405 410 415
 Leu Gly Ala Thr Glu Val Lys Leu Leu Gly His Gly Gln Pro Leu Asn
 420 425 430
 Trp Ile Ser Leu Glu Gln Asn Gly Ile Met Val Glu Leu Pro Gln Leu
 435 440 445
 Thr Ile His Gln Met Pro Cys Lys Trp Gly Trp Ala Leu Ala Leu Thr
 450 455 460

Asn Val Ile
 465

<210> 196
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 196
 tggtttgacc aggccaagtt cgg

23

<210> 197
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 197
 ggattcatcc tcaaggaaga gcgg

24

<210> 198
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 198
 aacttgacgc atcagccact ctgc

24

<210> 199
 <211> 45
 <212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 199

ttccgtgccc agcttcggta gcgagtgggt ctggtggtat tggca

45

<210> 200

<211> 2372

<212> DNA

<213> Homo sapiens

<400> 200

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catctgaggt gtttccctgg ctctgaaggg gtaggcacga tggccagggtg cttcagcctg 180
gtgttgcttc tcacttccat ctggaccacg aggctcctgg tccaaggctc tttgctgca 240
gaagagcttt ccatccaggt gtcatgcaga attatgggga tcacccttgt gagcaaaaag 300
gcgaaccagc agctgaattt cacagaagct aaggaggcct gtaggctgct gggactaagt 360
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aaaaatggg tgggtgtcct gatttggaag gttccagtga gccgacagtt tgcagcctat 540
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gccaatgata gcaaccctaa tgaggaatca aagaaaactg ataaaaacc agaagagtcc 1080
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tgtgacaaaa aattaaagca tttagaaaac tt 2372
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<210> 201

<211> 322
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic protein

<400> 201

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| 1 | | | | 5 | | | | 10 | | | | | | 15 | | |
| Thr | Arg | Leu | Leu | Val | Gln | Gly | Ser | Leu | Arg | Ala | Glu | Glu | Leu | Ser | Ile | |
| | | 20 | | | | | | 25 | | | | | 30 | | | |
| Gln | Val | Ser | Cys | Arg | Ile | Met | Gly | Ile | Thr | Leu | Val | Ser | Lys | Lys | Ala | |
| | | 35 | | | | | 40 | | | | | 45 | | | | |
| Asn | Gln | Gln | Leu | Asn | Phe | Thr | Glu | Ala | Lys | Glu | Ala | Cys | Arg | Leu | Leu | |
| | 50 | | | | | 55 | | | | | 60 | | | | | |
| Gly | Leu | Ser | Leu | Ala | Gly | Lys | Asp | Gln | Val | Glu | Thr | Ala | Leu | Lys | Ala | |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 | |
| Ser | Phe | Glu | Thr | Cys | Ser | Tyr | Gly | Trp | Val | Gly | Asp | Gly | Phe | Val | Val | |
| | | | | 85 | | | | | 90 | | | | | 95 | | |
| Ile | Ser | Arg | Ile | Ser | Pro | Asn | Pro | Lys | Cys | Gly | Lys | Asn | Gly | Val | Gly | |
| | | 100 | | | | | | 105 | | | | | 110 | | | |
| Val | Leu | Ile | Trp | Lys | Val | Pro | Val | Ser | Arg | Gln | Phe | Ala | Ala | Tyr | Cys | |
| | 115 | | | | | | 120 | | | | | 125 | | | | |
| Tyr | Asn | Ser | Ser | Asp | Thr | Trp | Thr | Asn | Ser | Cys | Ile | Pro | Glu | Ile | Ile | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Thr | Thr | Lys | Asp | Pro | Ile | Phe | Asn | Thr | Gln | Thr | Ala | Thr | Gln | Thr | Thr | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | |
| Glu | Phe | Ile | Val | Ser | Asp | Ser | Thr | Tyr | Ser | Val | Ala | Ser | Pro | Tyr | Ser | |
| | | | | 165 | | | | | 170 | | | | | 175 | | |
| Thr | Ile | Pro | Ala | Pro | Thr | Thr | Thr | Pro | Pro | Ala | Pro | Ala | Ser | Thr | Ser | |
| | | 180 | | | | | | 185 | | | | | 190 | | | |
| Ile | Pro | Arg | Arg | Lys | Lys | Leu | Ile | Cys | Val | Thr | Glu | Val | Phe | Met | Glu | |
| | 195 | | | | | 200 | | | | | | 205 | | | | |
| Thr | Ser | Thr | Met | Ser | Thr | Glu | Thr | Glu | Pro | Phe | Val | Glu | Asn | Lys | Ala | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Ala | Phe | Lys | Asn | Glu | Ala | Ala | Gly | Phe | Gly | Gly | Val | Pro | Thr | Ala | Leu | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | |
| Leu | Val | Leu | Ala | Leu | Leu | Phe | Phe | Gly | Ala | Ala | Ala | Gly | Leu | Gly | Phe | |
| | | | | 245 | | | | 250 | | | | | | 255 | | |
| Cys | Tyr | Val | Lys | Arg | Tyr | Val | Lys | Ala | Phe | Pro | Phe | Thr | Asn | Lys | Asn | |

260 265 270
 Gln Gln Lys Glu Met Ile Glu Thr Lys Val Val Lys Glu Glu Lys Ala
 275 280 285
 Asn Asp Ser Asn Pro Asn Glu Glu Ser Lys Lys Thr Asp Lys Asn Pro
 290 295 300
 Glu Glu Ser Lys Ser Pro Ser Lys Thr Thr Val Arg Cys Leu Glu Ala
 305 310 315 320
 Glu Val

<210> 202
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 202
 gagctttcca tccaggtgtc atgc 24

<210> 203
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 203
 gtcagtgaca gtacctactc gg 22

<210> 204
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 204
 tggagcagga ggagtagtag tagg 24

<210> 205
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic

oligonucleotide probe

<400> 205
aggaggcctg taggctgctg ggactaagtt tggccggcaa ggaccaagtt 50

<210> 206
<211> 1620
<212> DNA
<213> Homo sapiens

<220>
<221> modified_base
<222> (973)..(973)
<223> a, t, c or g

<220>
<221> modified_base
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<223> a, t, c or g

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<223> a, t, c or g

<220>
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<223> a, t, c or g

<400> 206
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actttccttt gtgtggtagg acttgaggaa gaaatccctt ggactttcac taaccctctg 1560

acatactccc cacaccagc tgaaggcttt ccgtaataaa aagattggga tttccttttg 1620

<210> 207

<211> 296

<212> PRT

<213> Homo sapiens

<400> 207

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Ser Ala Ala Phe Leu Leu Val Arg Lys Leu Pro Pro Leu Cys His Gly
35 40 45

Leu Pro Thr Gln Arg Glu Asp Gly Asn Pro Cys Asp Phe Asp Trp Arg
50 55 60

Glu Val Glu Ile Leu Met Phe Leu Ser Ala Ile Val Met Met Lys Asn
65 70 75 80

Arg Arg Ser Ile Thr Val Glu Gln His Ile Gly Asn Ile Phe Met Phe
85 90 95

Ser Lys Val Ala Asn Thr Ile Leu Phe Phe Arg Leu Asp Ile Arg Met
100 105 110

Gly Leu Leu Tyr Ile Thr Leu Cys Ile Val Phe Leu Met Thr Cys Lys
115 120 125

Pro Pro Leu Tyr Met Gly Pro Glu Tyr Ile Lys Tyr Phe Asn Asp Lys
130 135 140

Thr Ile Asp Glu Glu Leu Glu Arg Asp Lys Arg Val Thr Trp Ile Val
145 150 155 160

Glu Phe Phe Ala Asn Trp Ser Asn Asp Cys Gln Ser Phe Ala Pro Ile
165 170 175

Tyr Ala Asp Leu Ser Leu Lys Tyr Asn Cys Thr Gly Leu Asn Phe Gly
180 185 190

Lys Val Asp Val Gly Arg Tyr Thr Asp Val Ser Thr Arg Tyr Lys Val
195 200 205

Ser Thr Ser Pro Leu Thr Lys Gln Leu Pro Thr Leu Ile Leu Phe Gln
210 215 220

Gly Gly Lys Glu Ala Met Arg Arg Pro Gln Ile Asp Lys Lys Gly Arg
225 230 235 240

Ala Val Ser Trp Thr Phe Ser Glu Glu Asn Val Ile Arg Glu Phe Asn
245 250 255

Leu Asn Glu Leu Tyr Gln Arg Ala Lys Lys Leu Ser Lys Ala Gly Asp

| | | |
|---|-----|-----|
| 260 | 265 | 270 |
| Asn Ile Pro Glu Glu Gln Pro Val Ala Ser Thr Pro Thr Thr Val Ser | | |
| 275 | 280 | 285 |
| Asp Gly Glu Asn Lys Lys Asp Lys | | |
| 290 | 295 | |

<210> 208

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 208

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24

<210> 209

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 209

tggagacaat atccctgagg

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<210> 210

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 210

aacagttggc cacagcatgg cagg

24

<210> 211

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 211

ccattgatga ggaactagaa cgggacaaga gggtcacttg gattgtggag

50

<210> 212

<211> 1985
 <212> DNA
 <213> Homo sapiens

<400> 212

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aaaaaa 1985
  
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<210> 213
 <211> 360
 <212> PRT
 <213> Homo sapiens

<400> 213

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Met Gly Leu Leu Leu Leu Val Pro Leu Leu Leu Leu Pro Gly Ser Tyr
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Gly Leu Pro Phe Tyr Asn Gly Phe Tyr Tyr Ser Asn Ser Ala Asn Asp
  20                      25                     30

Gln Asn Leu Gly Asn Gly His Gly Lys Asp Leu Leu Asn Gly Val Lys
  35                      40                     45

Leu Val Val Glu Thr Pro Glu Glu Thr Leu Phe Thr Tyr Gln Gly Ala
  50                      55                     60
  
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Ser Val Ile Leu Pro Cys Arg Tyr Arg Tyr Glu Pro Ala Leu Val Ser
 65 70 75 80
 Pro Arg Arg Val Arg Val Lys Trp Trp Lys Leu Ser Glu Asn Gly Ala
 85 90 95
 Pro Glu Lys Asp Val Leu Val Ala Ile Gly Leu Arg His Arg Ser Phe
 100 105 110
 Gly Asp Tyr Gln Gly Arg Val His Leu Arg Gln Asp Lys Glu His Asp
 115 120 125
 Val Ser Leu Glu Ile Gln Asp Leu Arg Leu Glu Asp Tyr Gly Arg Tyr
 130 135 140
 Arg Cys Glu Val Ile Asp Gly Leu Glu Asp Glu Ser Gly Leu Val Glu
 145 150 155 160
 Leu Glu Leu Arg Gly Val Val Phe Pro Tyr Gln Ser Pro Asn Gly Arg
 165 170 175
 Tyr Gln Phe Asn Phe His Glu Gly Gln Gln Val Cys Ala Glu Gln Ala
 180 185 190
 Ala Val Val Ala Ser Phe Glu Gln Leu Phe Arg Ala Trp Glu Glu Gly
 195 200 205
 Leu Asp Trp Cys Asn Ala Gly Trp Leu Gln Asp Ala Thr Val Gln Tyr
 210 215 220
 Pro Ile Met Leu Pro Arg Gln Pro Cys Gly Gly Pro Gly Leu Ala Pro
 225 230 235 240
 Gly Val Arg Ser Tyr Gly Pro Arg His Arg Arg Leu His Arg Tyr Asp
 245 250 255
 Val Phe Cys Phe Ala Thr Ala Leu Lys Gly Arg Val Tyr Tyr Leu Glu
 260 265 270
 His Pro Glu Lys Leu Thr Leu Thr Glu Ala Arg Glu Ala Cys Gln Glu
 275 280 285
 Asp Asp Ala Thr Ile Ala Lys Val Gly Gln Leu Phe Ala Ala Trp Lys
 290 295 300
 Phe His Gly Leu Asp Arg Cys Asp Ala Gly Trp Leu Ala Asp Gly Ser
 305 310 315 320
 Val Arg Tyr Pro Val Val His Pro His Pro Asn Cys Gly Pro Pro Glu
 325 330 335
 Pro Gly Val Arg Ser Phe Gly Phe Pro Asp Pro Gln Ser Arg Leu Tyr
 340 345 350
 Gly Val Tyr Cys Tyr Arg Gln His
 355 360

<210> 214
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 214
 tgcttcgcta ctgccctc 18

 <210> 215
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 215
 ttcccttggtg ggttgag 18

 <210> 216
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 216
 agggctggaa gccagttc 18

 <210> 217
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

 <400> 217
 agccagtgtg gaaatgcg 18

 <210> 218
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 218
tgtccaaagt acacacacct gagg 24

<210> 219
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 219
gatgccacga tcgccaaggt gggacagctc tttgccgcct ggaag 45

<210> 220
<211> 1503
<212> DNA
<213> Homo sapiens

<400> 220
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aaa 1503

<210> 221
<211> 328
<212> PRT
<213> Homo sapiens

<400> 221
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1 5 10 15

| | | | | | | | | | | | | | | | | | | |
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| Gly | Ala | Gln | Gly | Lys | Pro | Ser | Pro | Asp | Ala | Gly | Pro | His | Gly | Gln | Gly | 20 | 25 | 30 |
| Arg | Val | His | Gln | Ala | Ala | Pro | Leu | Ser | Asp | Ala | Pro | His | Asp | Asp | Ala | 35 | 40 | 45 |
| His | Gly | Asn | Phe | Gln | Tyr | Asp | His | Glu | Ala | Phe | Leu | Gly | Arg | Glu | Val | 50 | 55 | 60 |
| Ala | Lys | Glu | Phe | Asp | Gln | Leu | Thr | Pro | Glu | Glu | Ser | Gln | Ala | Arg | Leu | 65 | 70 | 75 |
| Gly | Arg | Ile | Val | Asp | Arg | Met | Asp | Arg | Ala | Gly | Asp | Gly | Asp | Gly | Trp | 85 | 90 | 95 |
| Val | Ser | Leu | Ala | Glu | Leu | Arg | Ala | Trp | Ile | Ala | His | Thr | Gln | Gln | Arg | 100 | 105 | 110 |
| His | Ile | Arg | Asp | Ser | Val | Ser | Ala | Ala | Trp | Asp | Thr | Tyr | Asp | Thr | Asp | 115 | 120 | 125 |
| Arg | Asp | Gly | Arg | Val | Gly | Trp | Glu | Glu | Leu | Arg | Asn | Ala | Thr | Tyr | Gly | 130 | 135 | 140 |
| His | Tyr | Ala | Pro | Gly | Glu | Glu | Phe | His | Asp | Val | Glu | Asp | Ala | Glu | Thr | 145 | 150 | 155 |
| Tyr | Lys | Lys | Met | Leu | Ala | Arg | Asp | Glu | Arg | Arg | Phe | Arg | Val | Ala | Asp | 165 | 170 | 175 |
| Gln | Asp | Gly | Asp | Ser | Met | Ala | Thr | Arg | Glu | Glu | Leu | Thr | Ala | Phe | Leu | 180 | 185 | 190 |
| His | Pro | Glu | Glu | Phe | Pro | His | Met | Arg | Asp | Ile | Val | Ile | Ala | Glu | Thr | 195 | 200 | 205 |
| Leu | Glu | Asp | Leu | Asp | Arg | Asn | Lys | Asp | Gly | Tyr | Val | Gln | Val | Glu | Glu | 210 | 215 | 220 |
| Tyr | Ile | Ala | Asp | Leu | Tyr | Ser | Ala | Glu | Pro | Gly | Glu | Glu | Glu | Pro | Ala | 225 | 230 | 235 |
| Trp | Val | Gln | Thr | Glu | Arg | Gln | Gln | Phe | Arg | Asp | Phe | Arg | Asp | Leu | Asn | 245 | 250 | 255 |
| Lys | Asp | Gly | His | Leu | Asp | Gly | Ser | Glu | Val | Gly | His | Trp | Val | Leu | Pro | 260 | 265 | 270 |
| Pro | Ala | Gln | Asp | Gln | Pro | Leu | Val | Glu | Ala | Asn | His | Leu | Leu | His | Glu | 275 | 280 | 285 |
| Ser | Asp | Thr | Asp | Lys | Asp | Gly | Arg | Leu | Ser | Lys | Ala | Glu | Ile | Leu | Gly | 290 | 295 | 300 |
| Asn | Trp | Asn | Met | Phe | Val | Gly | Ser | Gln | Ala | Thr | Asn | Tyr | Gly | Glu | Asp | 305 | 310 | 315 |
| | | | | | | | | | | | | | | | | | | 320 |

Leu Thr Arg His His Asp Glu Leu
325

<210> 222

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 222

cgcaggccct catggccagg

20

<210> 223

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 223

gaaatcctgg gtaattgg

18

<210> 224

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 224

gtgcgcggtg ctcacagctc atc

23

<210> 225

<211> 44

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 225

ccccctgag cgacgctccc ccatgatgac gccacggga actt

44

<210> 226

<211> 2403

<212> DNA

<213> Homo sapiens

<400> 226

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aaa 2403
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<210> 227

<211> 550

<212> PRT

<213> Homo sapiens

<400> 227

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Leu Leu Pro Gly Pro Ala Gly Ser Glu Gly Ala Ala Pro Ile Ala Ile
          20             25             30

Thr Cys Phe Thr Arg Gly Leu Asp Ile Arg Lys Glu Lys Ala Asp Val
          35             40             45
```

| | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Leu | Cys | Pro | Gly | Gly | Cys | Pro | Leu | Glu | Glu | Phe | Ser | Val | Tyr | Gly | Asn | |
| 50 | | | | | | 55 | | | | | 60 | | | | | |
| Ile | Val | Tyr | Ala | Ser | Val | Ser | Ser | Ile | Cys | Gly | Ala | Ala | Val | His | Arg | |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 | |
| Gly | Val | Ile | Ser | Asn | Ser | Gly | Gly | Pro | Val | Arg | Val | Tyr | Ser | Leu | Pro | |
| | | | | 85 | | | | | 90 | | | | | 95 | | |
| Gly | Arg | Glu | Asn | Tyr | Ser | Ser | Val | Asp | Ala | Asn | Gly | Ile | Gln | Ser | Gln | |
| | | | 100 | | | | | 105 | | | | | 110 | | | |
| Met | Leu | Ser | Arg | Trp | Ser | Ala | Ser | Phe | Thr | Val | Thr | Lys | Gly | Lys | Ser | |
| | 115 | | | | | | 120 | | | | | 125 | | | | |
| Ser | Thr | Gln | Glu | Ala | Thr | Gly | Gln | Ala | Val | Ser | Thr | Ala | His | Pro | Pro | |
| | 130 | | | | | 135 | | | | | 140 | | | | | |
| Thr | Gly | Lys | Arg | Leu | Lys | Lys | Thr | Pro | Glu | Lys | Lys | Thr | Gly | Asn | Lys | |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 | |
| Asp | Cys | Lys | Ala | Asp | Ile | Ala | Phe | Leu | Ile | Asp | Gly | Ser | Phe | Asn | Ile | |
| | | | | 165 | | | | | 170 | | | | | 175 | | |
| Gly | Gln | Arg | Arg | Phe | Asn | Leu | Gln | Lys | Asn | Phe | Val | Gly | Lys | Val | Ala | |
| | | | 180 | | | | | 185 | | | | | 190 | | | |
| Leu | Met | Leu | Gly | Ile | Gly | Thr | Glu | Gly | Pro | His | Val | Gly | Leu | Val | Gln | |
| | 195 | | | | | | 200 | | | | | 205 | | | | |
| Ala | Ser | Glu | His | Pro | Lys | Ile | Glu | Phe | Tyr | Leu | Lys | Asn | Phe | Thr | Ser | |
| | 210 | | | | | 215 | | | | | 220 | | | | | |
| Ala | Lys | Asp | Val | Leu | Phe | Ala | Ile | Lys | Glu | Val | Gly | Phe | Arg | Gly | Gly | |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 | |
| Asn | Ser | Asn | Thr | Gly | Lys | Ala | Leu | Lys | His | Thr | Ala | Gln | Lys | Phe | Phe | |
| | | | | 245 | | | | | 250 | | | | | 255 | | |
| Thr | Val | Asp | Ala | Gly | Val | Arg | Lys | Gly | Ile | Pro | Lys | Val | Val | Val | Val | |
| | | | 260 | | | | | 265 | | | | | 270 | | | |
| Phe | Ile | Asp | Gly | Trp | Pro | Ser | Asp | Asp | Ile | Glu | Glu | Ala | Gly | Ile | Val | |
| | | 275 | | | | | 280 | | | | | 285 | | | | |
| Ala | Arg | Glu | Phe | Gly | Val | Asn | Val | Phe | Ile | Val | Ser | Val | Ala | Lys | Pro | |
| | 290 | | | | | 295 | | | | | 300 | | | | | |
| Ile | Pro | Glu | Glu | Leu | Gly | Met | Val | Gln | Asp | Val | Thr | Phe | Val | Asp | Lys | |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 | |
| Ala | Val | Cys | Arg | Asn | Asn | Gly | Phe | Phe | Ser | Tyr | His | Met | Pro | Asn | Trp | |
| | | | | 325 | | | | | 330 | | | | | 335 | | |
| Phe | Gly | Thr | Thr | Lys | Tyr | Val | Lys | Pro | Leu | Val | Gln | Lys | Leu | Cys | Thr | |
| | | | 340 | | | | | 345 | | | | | 350 | | | |

His Glu Gln Met Met Cys Ser Lys Thr Cys Tyr Asn Ser Val Asn Ile
 355 360 365
 Ala Phe Leu Ile Asp Gly Ser Ser Ser Val Gly Asp Ser Asn Phe Arg
 370 375 380
 Leu Met Leu Glu Phe Val Ser Asn Ile Ala Lys Thr Phe Glu Ile Ser
 385 390 395 400
 Asp Ile Gly Ala Lys Ile Ala Ala Val Gln Phe Thr Tyr Asp Gln Arg
 405 410 415
 Thr Glu Phe Ser Phe Thr Asp Tyr Ser Thr Lys Glu Asn Val Leu Ala
 420 425 430
 Val Ile Arg Asn Ile Arg Tyr Met Ser Gly Gly Thr Ala Thr Gly Asp
 435 440 445
 Ala Ile Ser Phe Thr Val Arg Asn Val Phe Gly Pro Ile Arg Glu Ser
 450 455 460
 Pro Asn Lys Asn Phe Leu Val Ile Val Thr Asp Gly Gln Ser Tyr Asp
 465 470 475 480
 Asp Val Gln Gly Pro Ala Ala Ala Ala His Asp Ala Gly Ile Thr Ile
 485 490 495
 Phe Ser Val Gly Val Ala Trp Ala Pro Leu Asp Asp Leu Lys Asp Met
 500 505 510
 Ala Ser Lys Pro Lys Glu Ser His Ala Phe Phe Thr Arg Glu Phe Thr
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 Gly Leu Glu Pro Ile Val Ser Asp Val Ile Arg Gly Ile Cys Arg Asp
 530 535 540
 Phe Leu Glu Ser Gln Gln
 545 550

<210> 228

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 228

tggtctcgca caccgatc

18

<210> 229

<211> 18

<212> DNA

<213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 229
 ctgctgtcca caggggag 18

<210> 230
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 230
 ccttgaagca tactgctc 18

<210> 231
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 231
 gagatagcaa tttccgcc 18

<210> 232
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 232
 ttcctcaaga gggcagcc 18

<210> 233
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide probe

<400> 233
 cttggcacca atgtccgaga tttc 24

<210> 234
 <211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide probe

<400> 234

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45

<210> 235

<211> 2586

<212> DNA

<213> Homo sapiens

<400> 235

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<213> Homo sapiens

<400> 236

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| Met | Gln | Arg | Leu | Gly | Ala | Thr | Leu | Leu | Cys | Leu | Leu | Leu | Ala | Ala | Ala | | | |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | | | | |
| Val | Pro | Thr | Ala | Pro | Ala | Pro | Ala | Pro | Thr | Ala | Thr | Ser | Ala | Pro | Val | | | |
| | | | 20 | | | | | 25 | | | | | 30 | | | | | |
| Lys | Pro | Gly | Pro | Ala | Leu | Ser | Tyr | Pro | Gln | Glu | Glu | Ala | Thr | Leu | Asn | | | |
| | | 35 | | | | | 40 | | | | | 45 | | | | | | |
| Glu | Met | Phe | Arg | Glu | Val | Glu | Glu | Leu | Met | Glu | Asp | Thr | Gln | His | Lys | | | |
| | 50 | | | | | 55 | | | | | 60 | | | | | | | |
| Leu | Arg | Ser | Ala | Val | Glu | Glu | Met | Glu | Ala | Glu | Glu | Ala | Ala | Ala | Lys | | | |
| 65 | | | | | 70 | | | | 75 | | | | | | 80 | | | |
| Ala | Ser | Ser | Glu | Val | Asn | Leu | Ala | Asn | Leu | Pro | Pro | Ser | Tyr | His | Asn | | | |
| | | | | 85 | | | | | 90 | | | | | 95 | | | | |
| Glu | Thr | Asn | Thr | Asp | Thr | Lys | Val | Gly | Asn | Asn | Thr | Ile | His | Val | His | | | |
| | | 100 | | | | | | 105 | | | | | 110 | | | | | |
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| Ser | Glu | Thr | Val | Ile | Thr | Ser | Val | Gly | Asp | Glu | Glu | Gly | Arg | Arg | Ser | | | |
| | 130 | | | | | 135 | | | | | 140 | | | | | | | |
| His | Glu | Cys | Ile | Ile | Asp | Glu | Asp | Cys | Gly | Pro | Ser | Met | Tyr | Cys | Gln | | | |
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| Phe | Ala | Ser | Phe | Gln | Tyr | Thr | Cys | Gln | Pro | Cys | Arg | Gly | Gln | Arg | Met | | | |
| | | | 165 | | | | | 170 | | | | | 175 | | | | | |
| Leu | Cys | Thr | Arg | Asp | Ser | Glu | Cys | Cys | Gly | Asp | Gln | Leu | Cys | Val | Trp | | | |
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| Gly | His | Cys | Thr | Lys | Met | Ala | Thr | Arg | Gly | Ser | Asn | Gly | Thr | Ile | Cys | | | |
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| Gly | Leu | Leu | Phe | Pro | Val | Cys | Thr | Pro | Leu | Pro | Val | Glu | Gly | Glu | Leu | | | |
| 225 | | | | | 230 | | | | 235 | | | | | 240 | | | | |
| Cys | His | Asp | Pro | Ala | Ser | Arg | Leu | Leu | Asp | Leu | Ile | Thr | Trp | Glu | Leu | | | |
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Glu Pro Asp Gly Ala Leu Asp Arg Cys Pro Cys Ala Ser Gly Leu Leu
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Cys Gln Pro His Ser His Ser Leu Val Tyr Val Cys Lys Pro Thr Phe
275 280 285

Val Gly Ser Arg Asp Gln Asp Gly Glu Ile Leu Leu Pro Arg Glu Val
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24

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<212> PRT

<213> Homo Sapien

<400> 245

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| Met | Arg | Leu | Leu | Val | Ala | Pro | Leu | Leu | Leu | Ala | Trp | Val | Ala | Gly | 1 | 5 | 10 | 15 |
| Ala | Thr | Ala | Thr | Val | Pro | Val | Val | Pro | Trp | His | Val | Pro | Cys | Pro | 20 | 25 | 30 | |
| Pro | Gln | Cys | Ala | Cys | Gln | Ile | Arg | Pro | Trp | Tyr | Thr | Pro | Arg | Ser | 35 | 40 | 45 | |
| Ser | Tyr | Arg | Glu | Ala | Thr | Thr | Val | Asp | Cys | Asn | Asp | Leu | Phe | Leu | 50 | 55 | 60 | |
| Thr | Ala | Val | Pro | Pro | Ala | Leu | Pro | Ala | Gly | Thr | Gln | Thr | Leu | Leu | 65 | 70 | 75 | |
| Leu | Gln | Ser | Asn | Ser | Ile | Val | Arg | Val | Asp | Gln | Ser | Glu | Leu | Gly | 80 | 85 | 90 | |
| Tyr | Leu | Ala | Asn | Leu | Thr | Glu | Leu | Asp | Leu | Ser | Gln | Asn | Ser | Phe | 95 | 100 | 105 | |
| Ser | Asp | Ala | Arg | Asp | Cys | Asp | Phe | His | Ala | Leu | Pro | Gln | Leu | Leu | 110 | 115 | 120 | |
| Ser | Leu | His | Leu | Glu | Glu | Asn | Gln | Leu | Thr | Arg | Leu | Glu | Asp | His | 125 | 130 | 135 | |
| Ser | Phe | Ala | Gly | Leu | Ala | Ser | Leu | Gln | Glu | Leu | Tyr | Leu | Asn | His | 140 | 145 | 150 | |

| | | |
|---------------------|-----------------|-------------------------|
| Asn Gln Leu Tyr Arg | Ile Ala Pro Arg | Ala Phe Ser Gly Leu Ser |
| 155 | 160 | 165 |
| Asn Leu Leu Arg Leu | His Leu Asn Ser | Asn Leu Leu Arg Ala Ile |
| 170 | 175 | 180 |
| Asp Ser Arg Trp Phe | Glu Met Leu Pro | Asn Leu Glu Ile Leu Met |
| 185 | 190 | 195 |
| Ile Gly Gly Asn Lys | Val Asp Ala Ile | Leu Asp Met Asn Phe Arg |
| 200 | 205 | 210 |
| Pro Leu Ala Asn Leu | Arg Ser Leu Val | Leu Ala Gly Met Asn Leu |
| 215 | 220 | 225 |
| Arg Glu Ile Ser Asp | Tyr Ala Leu Glu | Gly Leu Gln Ser Leu Glu |
| 230 | 235 | 240 |
| Ser Leu Ser Phe Tyr | Asp Asn Gln Leu | Ala Arg Val Pro Arg Arg |
| 245 | 250 | 255 |
| Ala Leu Glu Gln Val | Pro Gly Leu Lys | Phe Leu Asp Leu Asn Lys |
| 260 | 265 | 270 |
| Asn Pro Leu Gln Arg | Val Gly Pro Gly | Asp Phe Ala Asn Met Leu |
| 275 | 280 | 285 |
| His Leu Lys Glu Leu | Gly Leu Asn Asn | Met Glu Glu Leu Val Ser |
| 290 | 295 | 300 |
| Ile Asp Lys Phe Ala | Leu Val Asn Leu | Pro Glu Leu Thr Lys Leu |
| 305 | 310 | 315 |
| Asp Ile Thr Asn Asn | Pro Arg Leu Ser | Phe Ile His Pro Arg Ala |
| 320 | 325 | 330 |
| Phe His His Leu Pro | Gln Met Glu Thr | Leu Met Leu Asn Asn Asn |
| 335 | 340 | 345 |
| Ala Leu Ser Ala Leu | His Gln Gln Thr | Val Glu Ser Leu Pro Asn |
| 350 | 355 | 360 |
| Leu Gln Glu Val Gly | Leu His Gly Asn | Pro Ile Arg Cys Asp Cys |
| 365 | 370 | 375 |
| Val Ile Arg Trp Ala | Asn Ala Thr Gly | Thr Arg Val Arg Phe Ile |
| 380 | 385 | 390 |
| Glu Pro Gln Ser Thr | Leu Cys Ala Glu | Pro Pro Asp Leu Gln Arg |
| 395 | 400 | 405 |
| Leu Pro Val Arg Glu | Val Pro Phe Arg | Glu Met Thr Asp His Cys |
| 410 | 415 | 420 |
| Leu Pro Leu Ile Ser | Pro Arg Ser Phe | Pro Pro Ser Leu Gln Val |
| 425 | 430 | 435 |

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Ser | Gly | Glu | Ser | Met | Val | Leu | His | Cys | Arg | Ala | Leu | Ala | Glu | 440 | 445 | 450 |
| Pro | Glu | Pro | Glu | Ile | Tyr | Trp | Val | Thr | Pro | Ala | Gly | Leu | Arg | Leu | 455 | 460 | 465 |
| Thr | Pro | Ala | His | Ala | Gly | Arg | Arg | Tyr | Arg | Val | Tyr | Pro | Glu | Gly | 470 | 475 | 480 |
| Thr | Leu | Glu | Leu | Arg | Arg | Val | Thr | Ala | Glu | Glu | Ala | Gly | Leu | Tyr | 485 | 490 | 495 |
| Thr | Cys | Val | Ala | Gln | Asn | Leu | Val | Gly | Ala | Asp | Thr | Lys | Thr | Val | 500 | 505 | 510 |
| Ser | Val | Val | Val | Gly | Arg | Ala | Leu | Leu | Gln | Pro | Gly | Arg | Asp | Glu | 515 | 520 | 525 |
| Gly | Gln | Gly | Leu | Glu | Leu | Arg | Val | Gln | Glu | Thr | His | Pro | Tyr | His | 530 | 535 | 540 |
| Ile | Leu | Leu | Ser | Trp | Val | Thr | Pro | Pro | Asn | Thr | Val | Ser | Thr | Asn | 545 | 550 | 555 |
| Leu | Thr | Trp | Ser | Ser | Ala | Ser | Ser | Leu | Arg | Gly | Gln | Gly | Ala | Thr | 560 | 565 | 570 |
| Ala | Leu | Ala | Arg | Leu | Pro | Arg | Gly | Thr | His | Ser | Tyr | Asn | Ile | Thr | 575 | 580 | 585 |
| Arg | Leu | Leu | Gln | Ala | Thr | Glu | Tyr | Trp | Ala | Cys | Leu | Gln | Val | Ala | 590 | 595 | 600 |
| Phe | Ala | Asp | Ala | His | Thr | Gln | Leu | Ala | Cys | Val | Trp | Ala | Arg | Thr | 605 | 610 | 615 |
| Lys | Glu | Ala | Thr | Ser | Cys | His | Arg | Ala | Leu | Gly | Asp | Arg | Pro | Gly | 620 | 625 | 630 |
| Leu | Ile | Ala | Ile | Leu | Ala | Leu | Ala | Val | Leu | Leu | Leu | Ala | Ala | Gly | 635 | 640 | 645 |
| Leu | Ala | Ala | His | Leu | Gly | Thr | Gly | Gln | Pro | Arg | Lys | Gly | Val | Gly | 650 | 655 | 660 |
| Gly | Arg | Arg | Pro | Leu | Pro | Pro | Ala | Trp | Ala | Phe | Trp | Gly | Trp | Ser | 665 | 670 | 675 |
| Ala | Pro | Ser | Val | Arg | Val | Val | Ser | Ala | Pro | Leu | Val | Leu | Pro | Trp | 680 | 685 | 690 |
| Asn | Pro | Gly | Arg | Lys | Leu | Pro | Arg | Ser | Ser | Glu | Gly | Glu | Thr | Leu | 695 | 700 | 705 |
| Leu | Pro | Pro | Leu | Ser | Gln | Asn | Ser | | | | | | | | 710 | | |

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a 3401

<210> 250

<211> 546

<212> PRT

<213> Homo Sapien

<400> 250

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Arg | Gln | Thr | Ile | Ile | Lys | Val | Ile | Lys | Phe | Ile | Leu | Ile | Ile |
| 1 | | | | 5 | | | | | 10 | | | | | 15 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Cys | Tyr | Thr | Val | Tyr | Tyr | Val | His | Asn | Ile | Lys | Phe | Asp | Val | Asp |
| | | | | 20 | | | | | 25 | | | | | 30 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Cys | Thr | Val | Asp | Ile | Glu | Ser | Leu | Thr | Gly | Tyr | Arg | Thr | Tyr | Arg |
| | | | | 35 | | | | | 40 | | | | | 45 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Cys | Ala | His | Pro | Leu | Ala | Thr | Leu | Phe | Lys | Ile | Leu | Ala | Ser | Phe |
| | | | | 50 | | | | | 55 | | | | | 60 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Tyr | Ile | Ser | Leu | Val | Ile | Phe | Tyr | Gly | Leu | Ile | Cys | Met | Tyr | Thr |
| | | | | 65 | | | | | 70 | | | | | 75 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Trp | Trp | Met | Leu | Arg | Arg | Ser | Leu | Lys | Lys | Tyr | Ser | Phe | Glu |
| | | | | 80 | | | | | 85 | | | | | 90 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Ile | Arg | Glu | Glu | Ser | Ser | Tyr | Ser | Asp | Ile | Pro | Asp | Val | Lys |
| | | | | 95 | | | | | 100 | | | | | 105 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Asp | Phe | Ala | Phe | Met | Leu | His | Leu | Ile | Asp | Gln | Tyr | Asp | Pro |
| | | | | 110 | | | | | 115 | | | | | 120 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Tyr | Ser | Lys | Arg | Phe | Ala | Val | Phe | Leu | Ser | Glu | Val | Ser | Glu |
| | | | | 125 | | | | | 130 | | | | | 135 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Lys | Leu | Arg | Gln | Leu | Asn | Leu | Asn | Asn | Glu | Trp | Thr | Leu | Asp |
| | | | | 140 | | | | | 145 | | | | | 150 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Lys | Leu | Arg | Gln | Arg | Leu | Thr | Lys | Asn | Ala | Gln | Asp | Lys | Leu | Glu |
| | | | | 155 | | | | | 160 | | | | | 165 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | His | Leu | Phe | Met | Leu | Ser | Gly | Ile | Pro | Asp | Thr | Val | Phe | Asp |
| | | | | 170 | | | | | 175 | | | | | 180 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Val | Glu | Leu | Glu | Val | Leu | Lys | Leu | Glu | Leu | Ile | Pro | Asp | Val |
| | | | | 185 | | | | | 190 | | | | | 195 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Thr | Ile | Pro | Pro | Ser | Ile | Ala | Gln | Leu | Thr | Gly | Leu | Lys | Glu | Leu |
| | | | | 200 | | | | | 205 | | | | | 210 |

Trp Leu Tyr His Thr Ala Ala Lys Ile Glu Ala Pro Ala Leu Ala

| | | |
|-------------------------------------|-------------------------|-----|
| 215 | 220 | 225 |
| Phe Leu Arg Glu Asn Leu Arg Ala Leu | His Ile Lys Phe Thr Asp | |
| 230 | 235 | 240 |
| Ile Lys Glu Ile Pro Leu Trp Ile Tyr | Ser Leu Lys Thr Leu Glu | |
| 245 | 250 | 255 |
| Glu Leu His Leu Thr Gly Asn Leu Ser | Ala Glu Asn Asn Arg Tyr | |
| 260 | 265 | 270 |
| Ile Val Ile Asp Gly Leu Arg Glu Leu | Lys Arg Leu Lys Val Leu | |
| 275 | 280 | 285 |
| Arg Leu Lys Ser Asn Leu Ser Lys Leu | Pro Gln Val Val Thr Asp | |
| 290 | 295 | 300 |
| Val Gly Val His Leu Gln Lys Leu Ser | Ile Asn Asn Glu Gly Thr | |
| 305 | 310 | 315 |
| Lys Leu Ile Val Leu Asn Ser Leu Lys | Lys Met Ala Asn Leu Thr | |
| 320 | 325 | 330 |
| Glu Leu Glu Leu Ile Arg Cys Asp Leu | Glu Arg Ile Pro His Ser | |
| 335 | 340 | 345 |
| Ile Phe Ser Leu His Asn Leu Gln Glu | Ile Asp Leu Lys Asp Asn | |
| 350 | 355 | 360 |
| Asn Leu Lys Thr Ile Glu Glu Ile Ile | Ser Phe Gln His Leu His | |
| 365 | 370 | 375 |
| Arg Leu Thr Cys Leu Lys Leu Trp Tyr | Asn His Ile Ala Tyr Ile | |
| 380 | 385 | 390 |
| Pro Ile Gln Ile Gly Asn Leu Thr Asn | Leu Glu Arg Leu Tyr Leu | |
| 395 | 400 | 405 |
| Asn Arg Asn Lys Ile Glu Lys Ile Pro | Thr Gln Leu Phe Tyr Cys | |
| 410 | 415 | 420 |
| Arg Lys Leu Arg Tyr Leu Asp Leu Ser | His Asn Asn Leu Thr Phe | |
| 425 | 430 | 435 |
| Leu Pro Ala Asp Ile Gly Leu Leu Gln | Asn Leu Gln Asn Leu Ala | |
| 440 | 445 | 450 |
| Ile Thr Ala Asn Arg Ile Glu Thr Leu | Pro Pro Glu Leu Phe Gln | |
| 455 | 460 | 465 |
| Cys Arg Lys Leu Arg Ala Leu His Leu | Gly Asn Asn Val Leu Gln | |
| 470 | 475 | 480 |
| Ser Leu Pro Ser Arg Val Gly Glu Leu | Thr Asn Leu Thr Gln Ile | |
| 485 | 490 | 495 |

Glu Leu Arg Gly Asn Arg Leu Glu Cys Leu Pro Val Glu Leu Gly
500 505 510

Glu Cys Pro Leu Leu Lys Arg Ser Gly Leu Val Val Glu Glu Asp
515 520 525

Leu Phe Asn Thr Leu Pro Pro Glu Val Lys Glu Arg Leu Trp Arg
530 535 540

Ala Asp Lys Glu Gln Ala
545

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<213> Artificial Sequence

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<400> 252
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<210> 254
<211> 1650
<212> DNA
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<210> 255

<211> 452

<212> PRT

<213> Homo Sapien

<400> 255

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Met | Glu | Leu | Ala | Leu | Arg | Arg | Ser | Pro | Val | Pro | Arg | Trp | Leu | Leu | |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Leu | Leu | Pro | Leu | Leu | Leu | Gly | Leu | Asn | Ala | Gly | Ala | Val | Ile | Asp | |
| | | | | 20 | | | | | 25 | | | | | 30 | |
| Trp | Pro | Thr | Glu | Glu | Gly | Lys | Glu | Val | Trp | Asp | Tyr | Val | Thr | Val | |
| | | | | 35 | | | | | 40 | | | | | 45 | |
| Arg | Lys | Asp | Ala | Tyr | Met | Phe | Trp | Trp | Leu | Tyr | Tyr | Ala | Thr | Asn | |
| | | | | 50 | | | | | 55 | | | | | 60 | |
| Ser | Cys | Lys | Asn | Phe | Ser | Glu | Leu | Pro | Leu | Val | Met | Trp | Leu | Gln | |
| | | | | 65 | | | | | 70 | | | | | 75 | |
| Gly | Gly | Pro | Gly | Gly | Ser | Ser | Thr | Gly | Phe | Gly | Asn | Phe | Glu | Glu | |
| | | | | 80 | | | | | 85 | | | | | 90 | |
| Ile | Gly | Pro | Leu | Asp | Ser | Asp | Leu | Lys | Pro | Arg | Lys | Thr | Thr | Trp | |
| | | | | 95 | | | | | 100 | | | | | 105 | |
| Leu | Gln | Ala | Ala | Ser | Leu | Leu | Phe | Val | Asp | Asn | Pro | Val | Gly | Thr | |
| | | | | 110 | | | | | 115 | | | | | 120 | |
| Gly | Phe | Ser | Tyr | Val | Asn | Gly | Ser | Gly | Ala | Tyr | Ala | Lys | Asp | Leu | |
| | | | | 125 | | | | | 130 | | | | | 135 | |
| Ala | Met | Val | Ala | Ser | Asp | Met | Met | Val | Leu | Leu | Lys | Thr | Phe | Phe | |
| | | | | 140 | | | | | 145 | | | | | 150 | |
| Ser | Cys | His | Lys | Glu | Phe | Gln | Thr | Val | Pro | Phe | Tyr | Ile | Phe | Ser | |
| | | | | 155 | | | | | 160 | | | | | 165 | |
| Glu | Ser | Tyr | Gly | Gly | Lys | Met | Ala | Ala | Gly | Ile | Gly | Leu | Glu | Leu | |
| | | | | 170 | | | | | 175 | | | | | 180 | |
| Tyr | Lys | Ala | Ile | Gln | Arg | Gly | Thr | Ile | Lys | Cys | Asn | Phe | Ala | Gly | |
| | | | | 185 | | | | | 190 | | | | | 195 | |
| Val | Ala | Leu | Gly | Asp | Ser | Trp | Ile | Ser | Pro | Val | Asp | Ser | Val | Leu | |
| | | | | 200 | | | | | 205 | | | | | 210 | |
| Ser | Trp | Gly | Pro | Tyr | Leu | Tyr | Ser | Met | Ser | Leu | Leu | Glu | Asp | Lys | |
| | | | | 215 | | | | | 220 | | | | | 225 | |
| Gly | Leu | Ala | Glu | Val | Ser | Lys | Val | Ala | Glu | Gln | Val | Leu | Asn | Ala | |
| | | | | 230 | | | | | 235 | | | | | 240 | |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Val | Asn | Lys | Gly | Leu | Tyr | Arg | Glu | Ala | Thr | Glu | Leu | Trp | Gly | Lys | |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Ala | Glu | Met | Ile | Ile | Glu | Gln | Asn | Thr | Asp | Gly | Val | Asn | Phe | Tyr | |
| | | | | 260 | | | | | 265 | | | | | 270 | |
| Asn | Ile | Leu | Thr | Lys | Ser | Thr | Pro | Thr | Ser | Thr | Met | Glu | Ser | Ser | |
| | | | | 275 | | | | | 280 | | | | | 285 | |
| Leu | Glu | Phe | Thr | Gln | Ser | His | Leu | Val | Cys | Leu | Cys | Gln | Arg | His | |
| | | | | 290 | | | | | 295 | | | | | 300 | |
| Val | Arg | His | Leu | Gln | Arg | Asp | Ala | Leu | Ser | Gln | Leu | Met | Asn | Gly | |
| | | | | 305 | | | | | 310 | | | | | 315 | |
| Pro | Ile | Arg | Lys | Lys | Leu | Lys | Ile | Ile | Pro | Glu | Asp | Gln | Ser | Trp | |
| | | | | 320 | | | | | 325 | | | | | 330 | |
| Gly | Gly | Gln | Ala | Thr | Asn | Val | Phe | Val | Asn | Met | Glu | Glu | Asp | Phe | |
| | | | | 335 | | | | | 340 | | | | | 345 | |
| Met | Lys | Pro | Val | Ile | Ser | Ile | Val | Asp | Glu | Leu | Leu | Glu | Ala | Gly | |
| | | | | 350 | | | | | 355 | | | | | 360 | |
| Ile | Asn | Val | Thr | Val | Tyr | Asn | Gly | Gln | Leu | Asp | Leu | Ile | Val | Asp | |
| | | | | 365 | | | | | 370 | | | | | 375 | |
| Thr | Met | Gly | Gln | Glu | Ala | Trp | Val | Arg | Lys | Leu | Lys | Trp | Pro | Glu | |
| | | | | 380 | | | | | 385 | | | | | 390 | |
| Leu | Pro | Lys | Phe | Ser | Gln | Leu | Lys | Trp | Lys | Ala | Leu | Tyr | Ser | Asp | |
| | | | | 395 | | | | | 400 | | | | | 405 | |
| Pro | Lys | Ser | Leu | Glu | Thr | Ser | Ala | Phe | Val | Lys | Ser | Tyr | Lys | Asn | |
| | | | | 410 | | | | | 415 | | | | | 420 | |
| Leu | Ala | Phe | Tyr | Trp | Ile | Leu | Lys | Ala | Gly | His | Met | Val | Pro | Ser | |
| | | | | 425 | | | | | 430 | | | | | 435 | |
| Asp | Gln | Gly | Asp | Met | Ala | Leu | Lys | Met | Met | Arg | Leu | Val | Thr | Gln | |
| | | | | 440 | | | | | 445 | | | | | 450 | |

Gln Glu

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<211> 1100

<212> DNA

<213> Homo Sapien

<400> 256

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ccgttatcag gaccatgcgg ccgacgggtc atcacgtcgc gcatcgtggg 150

tggagaggac gccgaactcg ggcgttgcc gtggcagggg agcctgcgcc 200

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | 65 | | 70 | | 75 | | | | | | | | | |
| Trp | Ala | Leu | Thr | Ala | Ala | His | Cys | Phe | Glu | Thr | Tyr | Ser | Asp | Leu |
| | 80 | | | | | | | | 85 | | | | | 90 |
| Ser | Asp | Pro | Ser | Gly | Trp | Met | Val | Gln | Phe | Gly | Gln | Leu | Thr | Ser |
| | | | | 95 | | | | | 100 | | | | | 105 |
| Met | Pro | Ser | Phe | Trp | Ser | Leu | Gln | Ala | Tyr | Tyr | Thr | Arg | Tyr | Phe |
| | | | | 110 | | | | | 115 | | | | | 120 |
| Val | Ser | Asn | Ile | Tyr | Leu | Ser | Pro | Arg | Tyr | Leu | Gly | Asn | Ser | Pro |
| | | | | 125 | | | | | 130 | | | | | 135 |
| Tyr | Asp | Ile | Ala | Leu | Val | Lys | Leu | Ser | Ala | Pro | Val | Thr | Tyr | Thr |
| | | | | 140 | | | | | 145 | | | | | 150 |
| Lys | His | Ile | Gln | Pro | Ile | Cys | Leu | Gln | Ala | Ser | Thr | Phe | Glu | Phe |
| | | | | 155 | | | | | 160 | | | | | 165 |
| Glu | Asn | Arg | Thr | Asp | Cys | Trp | Val | Thr | Gly | Trp | Gly | Tyr | Ile | Lys |
| | | | | 170 | | | | | 175 | | | | | 180 |
| Glu | Asp | Glu | Ala | Leu | Pro | Ser | Pro | His | Thr | Leu | Gln | Glu | Val | Gln |
| | | | | 185 | | | | | 190 | | | | | 195 |
| Val | Ala | Ile | Ile | Asn | Asn | Ser | Met | Cys | Asn | His | Leu | Phe | Leu | Lys |
| | | | | 200 | | | | | 205 | | | | | 210 |
| Tyr | Ser | Phe | Arg | Lys | Asp | Ile | Phe | Gly | Asp | Met | Val | Cys | Ala | Gly |
| | | | | 215 | | | | | 220 | | | | | 225 |
| Asn | Ala | Gln | Gly | Gly | Lys | Asp | Ala | Cys | Phe | Gly | Asp | Ser | Gly | Gly |
| | | | | 230 | | | | | 235 | | | | | 240 |
| Pro | Leu | Ala | Cys | Asn | Lys | Asn | Gly | Leu | Trp | Tyr | Gln | Ile | Gly | Val |
| | | | | 245 | | | | | 250 | | | | | 255 |
| Val | Ser | Trp | Gly | Val | Gly | Cys | Gly | Arg | Pro | Asn | Arg | Pro | Gly | Val |
| | | | | 260 | | | | | 265 | | | | | 270 |
| Tyr | Thr | Asn | Ile | Ser | His | His | Phe | Glu | Trp | Ile | Gln | Lys | Leu | Met |
| | | | | 275 | | | | | 280 | | | | | 285 |
| Ala | Gln | Ser | Gly | Met | Ser | Gln | Pro | Asp | Pro | Ser | Trp | Pro | Leu | Leu |
| | | | | 290 | | | | | 295 | | | | | 300 |
| Phe | Phe | Pro | Leu | Leu | Trp | Ala | Leu | Pro | Leu | Leu | Gly | Pro | Val | |
| | | | | 305 | | | | | 310 | | | | | |

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<211> 2427

<212> DNA

<213> Homo Sapien

<400> 258

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cggagcccga ccagcggagg acgtgcccc caggctgggt gtccctgggc 150
cgtgaggacc ctgaggaaga gctgagtctc acctttgccc tgagacagca 200
gaatgtggaa agactctcgg agctgggtgca ggctgtgtcg gatcccagct 250
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 aatgattgat acctcaaatg taaaaaa 2427

<210> 259

<211> 556

<212> PRT

<213> Homo Sapien

<400> 259

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Gly | Leu | Gln | Ala | Cys | Leu | Leu | Gly | Leu | Phe | Ala | Leu | Ile | Leu |
| 1 | | | | 5 | | | | | 10 | | | | | 15 |
| Ser | Gly | Lys | Cys | Ser | Tyr | Ser | Pro | Glu | Pro | Asp | Gln | Arg | Arg | Thr |
| | | | | 20 | | | | | 25 | | | | | 30 |
| Leu | Pro | Pro | Gly | Trp | Val | Ser | Leu | Gly | Arg | Ala | Asp | Pro | Glu | Glu |
| | | | 35 | | | | | | 40 | | | | | 45 |
| Glu | Leu | Ser | Leu | Thr | Phe | Ala | Leu | Arg | Gln | Gln | Asn | Val | Glu | Arg |
| | | | 50 | | | | | | 55 | | | | | 60 |

| | | | |
|---|-----|-----|-----|
| Leu Ser Glu Leu Val Gln Ala Val Ser Asp Pro Ser Ser Pro Gln | 65 | 70 | 75 |
| Tyr Gly Lys Tyr Leu Thr Leu Glu Asn Val Ala Asp Leu Val Arg | 80 | 85 | 90 |
| Pro Ser Pro Leu Thr Leu His Thr Val Gln Lys Trp Leu Leu Ala | 95 | 100 | 105 |
| Ala Gly Ala Gln Lys Cys His Ser Val Ile Thr Gln Asp Phe Leu | 110 | 115 | 120 |
| Thr Cys Trp Leu Ser Ile Arg Gln Ala Glu Leu Leu Leu Pro Gly | 125 | 130 | 135 |
| Ala Glu Phe His His Tyr Val Gly Gly Pro Thr Glu Thr His Val | 140 | 145 | 150 |
| Val Arg Ser Pro His Pro Tyr Gln Leu Pro Gln Ala Leu Ala Pro | 155 | 160 | 165 |
| His Val Asp Phe Val Gly Gly Leu His Arg Phe Pro Pro Thr Ser | 170 | 175 | 180 |
| Ser Leu Arg Gln Arg Pro Glu Pro Gln Val Thr Gly Thr Val Gly | 185 | 190 | 195 |
| Leu His Leu Gly Val Thr Pro Ser Val Ile Arg Lys Arg Tyr Asn | 200 | 205 | 210 |
| Leu Thr Ser Gln Asp Val Gly Ser Gly Thr Ser Asn Asn Ser Gln | 215 | 220 | 225 |
| Ala Cys Ala Gln Phe Leu Glu Gln Tyr Phe His Asp Ser Asp Leu | 230 | 235 | 240 |
| Ala Gln Phe Met Arg Leu Phe Gly Gly Asn Phe Ala His Gln Ala | 245 | 250 | 255 |
| Ser Val Ala Arg Val Val Gly Gln Gln Gly Arg Gly Arg Ala Gly | 260 | 265 | 270 |
| Ile Glu Ala Ser Leu Asp Val Gln Tyr Leu Met Ser Ala Gly Ala | 275 | 280 | 285 |
| Asn Ile Ser Thr Trp Val Tyr Ser Ser Pro Gly Arg His Glu Gly | 290 | 295 | 300 |
| Gln Glu Pro Phe Leu Gln Trp Leu Met Leu Leu Ser Asn Glu Ser | 305 | 310 | 315 |
| Ala Leu Pro His Val His Thr Val Ser Tyr Gly Asp Asp Glu Asp | 320 | 325 | 330 |
| Ser Leu Ser Ser Ala Tyr Ile Gln Arg Val Asn Thr Glu Leu Met | 335 | 340 | 345 |

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Lys | Ala | Ala | Ala | Arg | Gly | Leu | Thr | Leu | Leu | Phe | Ala | Ser | Gly | Asp | 350 | 355 | 360 |
| Ser | Gly | Ala | Gly | Cys | Trp | Ser | Val | Ser | Gly | Arg | His | Gln | Phe | Arg | 365 | 370 | 375 |
| Pro | Thr | Phe | Pro | Ala | Ser | Ser | Pro | Tyr | Val | Thr | Thr | Val | Gly | Gly | 380 | 385 | 390 |
| Thr | Ser | Phe | Gln | Glu | Pro | Phe | Leu | Ile | Thr | Asn | Glu | Ile | Val | Asp | 395 | 400 | 405 |
| Tyr | Ile | Ser | Gly | Gly | Gly | Phe | Ser | Asn | Val | Phe | Pro | Arg | Pro | Ser | 410 | 415 | 420 |
| Tyr | Gln | Glu | Glu | Ala | Val | Thr | Lys | Phe | Leu | Ser | Ser | Ser | Pro | His | 425 | 430 | 435 |
| Leu | Pro | Pro | Ser | Ser | Tyr | Phe | Asn | Ala | Ser | Gly | Arg | Ala | Tyr | Pro | 440 | 445 | 450 |
| Asp | Val | Ala | Ala | Leu | Ser | Asp | Gly | Tyr | Trp | Val | Val | Ser | Asn | Arg | 455 | 460 | 465 |
| Val | Pro | Ile | Pro | Trp | Val | Ser | Gly | Thr | Ser | Ala | Ser | Thr | Pro | Val | 470 | 475 | 480 |
| Phe | Gly | Gly | Ile | Leu | Ser | Leu | Ile | Asn | Glu | His | Arg | Ile | Leu | Ser | 485 | 490 | 495 |
| Gly | Arg | Pro | Pro | Leu | Gly | Phe | Leu | Asn | Pro | Arg | Leu | Tyr | Gln | Gln | 500 | 505 | 510 |
| His | Gly | Ala | Gly | Leu | Phe | Asp | Val | Thr | Arg | Gly | Cys | His | Glu | Ser | 515 | 520 | 525 |
| Cys | Leu | Asp | Glu | Glu | Val | Glu | Gly | Gln | Gly | Phe | Cys | Ser | Gly | Pro | 530 | 535 | 540 |
| Gly | Trp | Asp | Pro | Val | Thr | Gly | Trp | Gly | Thr | Pro | Thr | Ser | Gln | Leu | 545 | 550 | 555 |

Cys

<210> 260

<211> 1638

<212> DNA

<213> Homo Sapien

<400> 260

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cgcgccccgg gcgggctgct cggcgcggaa cagtgctcgg catggcaggg 100

attccagggc tctcttctct tctcttcttt ctgctctgtg ctgttgggca 150

agtgagccct tacagtgtccc cctggaaacc cacttggcct gcataccgcc 200
tccctgtcgt cttgccccag tctaccctca atttagccaa gccagacttt 250
ggagccgaag ccaaattaga agtatcttct tcatgtggac cccagtgtca 300
taagggaact cactgtccca cttacgaaga ggccaagcaa tatctgtctt 350
atgaaacgct ctatgtccaat ggcagccgca cagagacgca ggtgggcatc 400
tacatcctca gcagtagtgg agatggggcc caacaccgag actcagggtc 450
ttcaggaaag tctcgaagga agcggcagat ttatggctat gacagcaggt 500
tcagcatttt tgggaaggac ttctgtctca actacccttt ctcaacatca 550
gtgaagttat ccacgggctg caccggcacc ctggtggcag agaagcatgt 600
cctcacagct gccactgca tacacgatgg aaaaacctat gtgaaaggaa 650
cccagaagct tcgagtgggc ttctaaagc ccaagtttaa agatgggtgg 700
cgaggggcca acgactccac ttcagccatg cccgagcaga tgaaatttca 750
gtggatccgg gtgaaacgca cccatgtgcc caagggttgg atcaagggca 800
atgccaatga catcggcatg gattatgatt atgccctcct ggaactcaaa 850
aagccccaca agagaaaatt tatgaagatt ggggtgagcc ctctgtctaa 900
gcagctgcca gggggcagaa ttcacttctc tggttatgac aatgaccgac 950
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ttggcatttt ttcagggcac cagtgggtgg acatgaatgg ttccccacag 1150
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ccaaattgtt ttttgtcatt ggcgtgcaca cgtgtgtgtg tgtgtgtgtg 1350
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caaactttga tttttatttc atctgaactt gtttcaaaga tttatatata 1600

atatttggca tacaagagat atgaaaaaaaa aaaaaaaa 1638

<210> 261

<211> 383

<212> PRT

<213> Homo Sapien

<400> 261

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Met | Ala | Gly | Ile | Pro | Gly | Leu | Leu | Phe | Leu | Leu | Phe | Phe | Leu | Leu | |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Cys | Ala | Val | Gly | Gln | Val | Ser | Pro | Tyr | Ser | Ala | Pro | Trp | Lys | Pro | |
| | | | | 20 | | | | | 25 | | | | | 30 | |
| Thr | Trp | Pro | Ala | Tyr | Arg | Leu | Pro | Val | Val | Leu | Pro | Gln | Ser | Thr | |
| | | | | 35 | | | | | 40 | | | | | 45 | |
| Leu | Asn | Leu | Ala | Lys | Pro | Asp | Phe | Gly | Ala | Glu | Ala | Lys | Leu | Glu | |
| | | | | 50 | | | | | 55 | | | | | 60 | |
| Val | Ser | Ser | Ser | Cys | Gly | Pro | Gln | Cys | His | Lys | Gly | Thr | Pro | Leu | |
| | | | | 65 | | | | | 70 | | | | | 75 | |
| Pro | Thr | Tyr | Glu | Glu | Ala | Lys | Gln | Tyr | Leu | Ser | Tyr | Glu | Thr | Leu | |
| | | | | 80 | | | | | 85 | | | | | 90 | |
| Tyr | Ala | Asn | Gly | Ser | Arg | Thr | Glu | Thr | Gln | Val | Gly | Ile | Tyr | Ile | |
| | | | | 95 | | | | | 100 | | | | | 105 | |
| Leu | Ser | Ser | Ser | Gly | Asp | Gly | Ala | Gln | His | Arg | Asp | Ser | Gly | Ser | |
| | | | | 110 | | | | | 115 | | | | | 120 | |
| Ser | Gly | Lys | Ser | Arg | Arg | Lys | Arg | Gln | Ile | Tyr | Gly | Tyr | Asp | Ser | |
| | | | | 125 | | | | | 130 | | | | | 135 | |
| Arg | Phe | Ser | Ile | Phe | Gly | Lys | Asp | Phe | Leu | Leu | Asn | Tyr | Pro | Phe | |
| | | | | 140 | | | | | 145 | | | | | 150 | |
| Ser | Thr | Ser | Val | Lys | Leu | Ser | Thr | Gly | Cys | Thr | Gly | Thr | Leu | Val | |
| | | | | 155 | | | | | 160 | | | | | 165 | |
| Ala | Glu | Lys | His | Val | Leu | Thr | Ala | Ala | His | Cys | Ile | His | Asp | Gly | |
| | | | | 170 | | | | | 175 | | | | | 180 | |
| Lys | Thr | Tyr | Val | Lys | Gly | Thr | Gln | Lys | Leu | Arg | Val | Gly | Phe | Leu | |
| | | | | 185 | | | | | 190 | | | | | 195 | |
| Lys | Pro | Lys | Phe | Lys | Asp | Gly | Gly | Arg | Gly | Ala | Asn | Asp | Ser | Thr | |
| | | | | 200 | | | | | 205 | | | | | 210 | |
| Ser | Ala | Met | Pro | Glu | Gln | Met | Lys | Phe | Gln | Trp | Ile | Arg | Val | Lys | |
| | | | | 215 | | | | | 220 | | | | | 225 | |
| Arg | Thr | His | Val | Pro | Lys | Gly | Trp | Ile | Lys | Gly | Asn | Ala | Asn | Asp | |
| | | | | 230 | | | | | 235 | | | | | 240 | |

| | | | | |
|-------------------------------------|-------------------------|-----|-----|-----|
| Ile Gly Met Asp Tyr Asp Tyr Ala Leu | Leu Glu Leu Lys Lys Pro | 245 | 250 | 255 |
| His Lys Arg Lys Phe Met Lys Ile Gly | Val Ser Pro Pro Ala Lys | 260 | 265 | 270 |
| Gln Leu Pro Gly Gly Arg Ile His Phe | Ser Gly Tyr Asp Asn Asp | 275 | 280 | 285 |
| Arg Pro Gly Asn Leu Val Tyr Arg Phe | Cys Asp Val Lys Asp Glu | 290 | 295 | 300 |
| Thr Tyr Asp Leu Leu Tyr Gln Gln Cys | Asp Ala Gln Pro Gly Ala | 305 | 310 | 315 |
| Ser Gly Ser Gly Val Tyr Val Arg Met | Trp Lys Arg Gln Gln Gln | 320 | 325 | 330 |
| Lys Trp Glu Arg Lys Ile Ile Gly Ile | Phe Ser Gly His Gln Trp | 335 | 340 | 345 |
| Val Asp Met Asn Gly Ser Pro Gln Asp | Phe Asn Val Ala Val Arg | 350 | 355 | 360 |
| Ile Thr Pro Leu Lys Tyr Ala Gln Ile | Cys Tyr Trp Ile Lys Gly | 365 | 370 | 375 |
| Asn Tyr Leu Asp Cys Arg Glu Gly | | 380 | | |

<210> 262
 <211> 1378
 <212> DNA
 <213> Homo Sapien

<400> 262
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 caggatacct gttccccag cctgtgggaa gccccagcag ctgaaccggg 200
 ttgtgggagg cgaggacagc actgacagcg agtggccctg gatcgtgagc 250
 atccagaaga atgggaccca ccaactgcgc ggttctctgc tcaccagccg 300
 ctgggtgatc actgctgccc actgtttcaa ggacaacctg aacaaaccat 350
 acctgttctc tgtgctgctg ggggcctggc agctggggaa ccctggctct 400
 cgggtcccaga aggtgggtgt tgcctgggtg gagccccacc ctgtgtattc 450
 ctggaaggaa ggtgcctgtg cagacattgc cctgggtgcgt ctcgagcgct 500

ccatacagtt ctcaagagcgg gtcttgccca tctgcctacc tgatgcctct 550
 atccacctcc ctccaaacac ccactgctgg atctcaggct gggggagcat 600
 ccaagatgga gttcccttgc cccaccctca gaccctgcag aagctgaagg 650
 ttcctatcat cgactcggaa gtctgcagcc atctgtactg gcggggagca 700
 ggacagggac ccactactga ggacatgctg tgtgccggct acttggaggg 750
 ggagcgggat gcttgtctgg gcgactccgg gggccccctc atgtgccagg 800
 tggacggcgc ctggctgctg gccggcatca tcagctgggg cgagggctgt 850
 gccgagcgca acaggcccgg ggtctacatc agcctctctg cgcaccgctc 900
 ctgggtggag aagatcgtgc aaggggtgca gctccgcggg cgcgctcagg 950
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 cacatctgga tctggatctg cggcggcctc gggcggtttc ccccgccgta 1100
 aataggctca tctaccteta cctctggggg cccggacggc tgctgcggaa 1150
 aggaaacccc ctccccgacc cgcccagcgg cctcaggccc ccctccaagg 1200
 catcaggccc cgcccaacgg cctcatgtcc ccgccccac gacttccggc 1250
 cccgcccccg ggccccagcg cttttgtgta tataaatgtt aatgattttt 1300
 ataggtattt gtaaccctgc ccacatatct tatttattcc tccaatttca 1350
 ataaattatt tattctccaa aaaaaaaaa 1378

<210> 263
 <211> 317
 <212> PRT
 <213> Homo Sapien

<400> 263

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Val | Val | Ser | Gly | Ala | Pro | Pro | Ala | Leu | Gly | Gly | Gly | Cys | Leu |
| 1 | | | | 5 | | | | | 10 | | | | | 15 |
| Gly | Thr | Phe | Thr | Ser | Leu | Leu | Leu | Leu | Ala | Ser | Thr | Ala | Ile | Leu |
| | | | | 20 | | | | | 25 | | | | | 30 |
| Asn | Ala | Ala | Arg | Ile | Pro | Val | Pro | Pro | Ala | Cys | Gly | Lys | Pro | Gln |
| | | | | 35 | | | | | 40 | | | | | 45 |
| Gln | Leu | Asn | Arg | Val | Val | Gly | Gly | Glu | Asp | Ser | Thr | Asp | Ser | Glu |
| | | | | 50 | | | | | 55 | | | | | 60 |
| Trp | Pro | Trp | Ile | Val | Ser | Ile | Gln | Lys | Asn | Gly | Thr | His | His | Cys |
| | | | | 65 | | | | | 70 | | | | | 75 |

| | | | |
|-----------------|---------------------|-------------------------|-----|
| Ala Gly Ser Leu | Leu Thr Ser Arg Trp | Val Ile Thr Ala Ala His | |
| | 80 | 85 | 90 |
| Cys Phe Lys Asp | Asn Leu Asn Lys Pro | Tyr Leu Phe Ser Val Leu | |
| | 95 | 100 | 105 |
| Leu Gly Ala Trp | Gln Leu Gly Asn Pro | Gly Ser Arg Ser Gln Lys | |
| | 110 | 115 | 120 |
| Val Gly Val Ala | Trp Val Glu Pro His | Pro Val Tyr Ser Trp Lys | |
| | 125 | 130 | 135 |
| Glu Gly Ala Cys | Ala Asp Ile Ala Leu | Val Arg Leu Glu Arg Ser | |
| | 140 | 145 | 150 |
| Ile Gln Phe Ser | Glu Arg Val Leu Pro | Ile Cys Leu Pro Asp Ala | |
| | 155 | 160 | 165 |
| Ser Ile His Leu | Pro Pro Asn Thr His | Cys Trp Ile Ser Gly Trp | |
| | 170 | 175 | 180 |
| Gly Ser Ile Gln | Asp Gly Val Pro Leu | Pro His Pro Gln Thr Leu | |
| | 185 | 190 | 195 |
| Gln Lys Leu Lys | Val Pro Ile Ile Asp | Ser Glu Val Cys Ser His | |
| | 200 | 205 | 210 |
| Leu Tyr Trp Arg | Gly Ala Gly Gln Gly | Pro Ile Thr Glu Asp Met | |
| | 215 | 220 | 225 |
| Leu Cys Ala Gly | Tyr Leu Glu Gly Glu | Arg Asp Ala Cys Leu Gly | |
| | 230 | 235 | 240 |
| Asp Ser Gly Gly | Pro Leu Met Cys Gln | Val Asp Gly Ala Trp Leu | |
| | 245 | 250 | 255 |
| Leu Ala Gly Ile | Ile Ser Trp Gly Glu | Gly Cys Ala Glu Arg Asn | |
| | 260 | 265 | 270 |
| Arg Pro Gly Val | Tyr Ile Ser Leu Ser | Ala His Arg Ser Trp Val | |
| | 275 | 280 | 285 |
| Glu Lys Ile Val | Gln Gly Val Gln Leu | Arg Gly Arg Ala Gln Gly | |
| | 290 | 295 | 300 |
| Gly Gly Ala Leu | Arg Ala Pro Ser Gln | Gly Ser Gly Ala Ala Ala | |
| | 305 | 310 | 315 |

Arg Ser

<210> 264

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 264

gtccgcaagg atgcctacat gttc 24

<210> 265

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 265

gcagaggtgt ctaaggttg 19

<210> 266

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 266

agctctagac caatgccagc ttcc 24

<210> 267

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 267

gccaccaact cctgcaagaa cttctcagaa ctgcccctgg tcatg 45

<210> 268

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 268

ggggaattca ccctatgaca ttgcc 25

<210> 269

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 269
gaatgccctg caagcatcaa ctgg 24

<210> 270
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 270
gcacctgtca cctacactaa acacatccag cccatctgtc tccaggcctc 50

<210> 271
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 271
gcggaagggc agaatgggac tccaag 26

<210> 272
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 272
cagccctgcc acatgtgc 18

<210> 273
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 273
tactgggtgg tcagcaac 18

<210> 274
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 274
ggcgaagagc aggggtgagac cccg 24

<210> 275

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 275

gccctcatcc tctctggcaa atgcagttac agcccggagc ccgac 45

<210> 276

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 276

gggcagggat tccagggctc c 21

<210> 277

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 277

ggctatgaca gcaggttc 18

<210> 278

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 278

tgacaatgac cgaccagg 18

<210> 279

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 279

gcatcgcatt gctggtagag caag 24

<210> 280
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 280
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<210> 281
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 281
cgtctcgagc gtcctatata gttcccttgc ccca 34

<210> 282
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 282
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tgccaggtgg a 61

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<211> 119
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

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atgctgtgtg ccggctact 119

<210> 284
<211> 1875
<212> DNA
<213> Homo Sapien

<400> 284
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aggtatcccc gacggcctca gacatgctgc acatgagatg ggacgaggag 200
ctggccgcct tcgccaaggc ctacgcacgg cagtgcgtgt ggggccacaa 250
caaggagcgc gggcgccgcg gcgagaatct gttcgccatc acagacgagg 300
gcatggacgt gccgctggcc atggaggagt ggccaccaga gcgtgagcac 350
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cacgcaggtg gtatgggcca agacagagag gatcggctgt ggttcccact 450
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<210> 285

<211> 463

<212> PRT

<213> Homo Sapien

<400> 285

| | | | | | | | | | | | | | | | | | | |
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| Met | His | Gly | Ser | Cys | Ser | Phe | Leu | Met | Leu | Leu | Leu | Pro | Leu | Leu | 1 | 5 | 10 | 15 |
| Leu | Leu | Leu | Val | Ala | Thr | Thr | Gly | Pro | Val | Gly | Ala | Leu | Thr | Asp | 20 | 25 | 30 | |
| Glu | Glu | Lys | Arg | Leu | Met | Val | Glu | Leu | His | Asn | Leu | Tyr | Arg | Ala | 35 | 40 | 45 | |
| Gln | Val | Ser | Pro | Thr | Ala | Ser | Asp | Met | Leu | His | Met | Arg | Trp | Asp | 50 | 55 | 60 | |
| Glu | Glu | Leu | Ala | Ala | Phe | Ala | Lys | Ala | Tyr | Ala | Arg | Gln | Cys | Val | 65 | 70 | 75 | |
| Trp | Gly | His | Asn | Lys | Glu | Arg | Gly | Arg | Arg | Gly | Glu | Asn | Leu | Phe | 80 | 85 | 90 | |
| Ala | Ile | Thr | Asp | Glu | Gly | Met | Asp | Val | Pro | Leu | Ala | Met | Glu | Glu | 95 | 100 | 105 | |
| Trp | His | His | Glu | Arg | Glu | His | Tyr | Asn | Leu | Ser | Ala | Ala | Thr | Cys | 110 | 115 | 120 | |
| Ser | Pro | Gly | Gln | Met | Cys | Gly | His | Tyr | Thr | Gln | Val | Val | Trp | Ala | 125 | 130 | 135 | |
| Lys | Thr | Glu | Arg | Ile | Gly | Cys | Gly | Ser | His | Phe | Cys | Glu | Lys | Leu | 140 | 145 | 150 | |
| Gln | Gly | Val | Glu | Glu | Thr | Asn | Ile | Glu | Leu | Leu | Val | Cys | Asn | Tyr | 155 | 160 | 165 | |

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|-----|-----|-----|
| Glu | Pro | Pro | Gly | Asn | Val | Lys | Gly | Lys | Arg | Pro | Tyr | Gln | Glu | Gly | | 170 | 175 | 180 |
| Thr | Pro | Cys | Ser | Gln | Cys | Pro | Ser | Gly | Tyr | His | Cys | Lys | Asn | Ser | | 185 | 190 | 195 |
| Leu | Cys | Glu | Pro | Ile | Gly | Ser | Pro | Glu | Asp | Ala | Gln | Asp | Leu | Pro | | 200 | 205 | 210 |
| Tyr | Leu | Val | Thr | Glu | Ala | Pro | Ser | Phe | Arg | Ala | Thr | Glu | Ala | Ser | | 215 | 220 | 225 |
| Asp | Ser | Arg | Lys | Met | Gly | Thr | Pro | Ser | Ser | Leu | Ala | Thr | Gly | Ile | | 230 | 235 | 240 |
| Pro | Ala | Phe | Leu | Val | Thr | Glu | Val | Ser | Gly | Ser | Leu | Ala | Thr | Lys | | 245 | 250 | 255 |
| Ala | Leu | Pro | Ala | Val | Glu | Thr | Gln | Ala | Pro | Thr | Ser | Leu | Ala | Thr | | 260 | 265 | 270 |
| Lys | Asp | Pro | Pro | Ser | Met | Ala | Thr | Glu | Ala | Pro | Pro | Cys | Val | Thr | | 275 | 280 | 285 |
| Thr | Glu | Val | Pro | Ser | Ile | Leu | Ala | Ala | His | Ser | Leu | Pro | Ser | Leu | | 290 | 295 | 300 |
| Asp | Glu | Glu | Pro | Val | Thr | Phe | Pro | Lys | Ser | Thr | His | Val | Pro | Ile | | 305 | 310 | 315 |
| Pro | Lys | Ser | Ala | Asp | Lys | Val | Thr | Asp | Lys | Thr | Lys | Val | Pro | Ser | | 320 | 325 | 330 |
| Arg | Ser | Pro | Glu | Asn | Ser | Leu | Asp | Pro | Lys | Met | Ser | Leu | Thr | Gly | | 335 | 340 | 345 |
| Ala | Arg | Glu | Leu | Leu | Pro | His | Ala | Gln | Glu | Glu | Ala | Glu | Ala | Glu | | 350 | 355 | 360 |
| Ala | Glu | Leu | Pro | Pro | Ser | Ser | Glu | Val | Leu | Ala | Ser | Val | Phe | Pro | | 365 | 370 | 375 |
| Ala | Gln | Asp | Lys | Pro | Gly | Glu | Leu | Gln | Ala | Thr | Leu | Asp | His | Thr | | 380 | 385 | 390 |
| Gly | His | Thr | Ser | Ser | Lys | Ser | Leu | Pro | Asn | Phe | Pro | Asn | Thr | Ser | | 395 | 400 | 405 |
| Ala | Thr | Ala | Asn | Ala | Thr | Gly | Gly | Arg | Ala | Leu | Ala | Leu | Gln | Ser | | 410 | 415 | 420 |
| Ser | Leu | Pro | Gly | Ala | Glu | Gly | Pro | Asp | Lys | Pro | Ser | Val | Val | Ser | | 425 | 430 | 435 |
| Gly | Leu | Asn | Ser | Gly | Pro | Gly | His | Val | Trp | Gly | Pro | Leu | Leu | Gly | | 440 | 445 | 450 |

Leu Leu Leu Leu Pro Pro Leu Val Leu Ala Gly Ile Phe
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<210> 286

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 286

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<210> 287

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 287

ctcatattgc acaccagtaa ttcg 24

<210> 288

<211> 45

<212> DNA

<213> Artificial Sequence

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<223> Synthetic Oligonucleotide Probe

<400> 288

atgaggagaa acgtttgatg gtggagctgc acaacctcta ccggg 45

<210> 289

<211> 3662

<212> DNA

<213> Homo Sapien

<400> 289

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 <212> PRT
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<400> 290
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 Ala Ser Ser Met Ser His Leu Gln Ser Leu Arg Glu Val Lys Leu
 35 40 45
 Asn Asn Asn Glu Leu Glu Thr Ile Pro Asn Leu Gly Pro Val Ser
 50 55 60
 Ala Asn Ile Thr Leu Leu Ser Leu Ala Gly Asn Arg Ile Val Glu
 65 70 75
 Ile Leu Pro Glu His Leu Lys Glu Phe Gln Ser Leu Glu Thr Leu
 80 85 90
 Asp Leu Ser Ser Asn Asn Ile Ser Glu Leu Gln Thr Ala Phe Pro
 95 100 105
 Ala Leu Gln Leu Lys Tyr Leu Tyr Leu Asn Ser Asn Arg Val Thr
 110 115 120
 Ser Met Glu Pro Gly Tyr Phe Asp Asn Leu Ala Asn Thr Leu Leu
 125 130 135
 Val Leu Lys Leu Asn Arg Asn Arg Ile Ser Ala Ile Pro Pro Lys
 140 145 150
 Met Phe Lys Leu Pro Gln Leu Gln His Leu Glu Leu Asn Arg Asn

| | | | | | |
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| | 155 | | 160 | | 165 |
| Lys Ile Lys Asn Val Asp Gly Leu Thr Phe Gln Gly Leu Gly Ala | 170 | | 175 | | 180 |
| Leu Lys Ser Leu Lys Met Gln Arg Asn Gly Val Thr Lys Leu Met | 185 | | 190 | | 195 |
| Asp Gly Ala Phe Trp Gly Leu Ser Asn Met Glu Ile Leu Gln Leu | 200 | | 205 | | 210 |
| Asp His Asn Asn Leu Thr Glu Ile Thr Lys Gly Trp Leu Tyr Gly | 215 | | 220 | | 225 |
| Leu Leu Met Leu Gln Glu Leu His Leu Ser Gln Asn Ala Ile Asn | 230 | | 235 | | 240 |
| Arg Ile Ser Pro Asp Ala Trp Glu Phe Cys Gln Lys Leu Ser Glu | 245 | | 250 | | 255 |
| Leu Asp Leu Thr Phe Asn His Leu Ser Arg Leu Asp Asp Ser Ser | 260 | | 265 | | 270 |
| Phe Leu Gly Leu Ser Leu Leu Asn Thr Leu His Ile Gly Asn Asn | 275 | | 280 | | 285 |
| Arg Val Ser Tyr Ile Ala Asp Cys Ala Phe Arg Gly Leu Ser Ser | 290 | | 295 | | 300 |
| Leu Lys Thr Leu Asp Leu Lys Asn Asn Glu Ile Ser Trp Thr Ile | 305 | | 310 | | 315 |
| Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Arg | 320 | | 325 | | 330 |
| Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Ala | 335 | | 340 | | 345 |
| Phe Thr Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp Asn | 350 | | 355 | | 360 |
| Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Gln Met Lys Lys | 365 | | 370 | | 375 |
| Leu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cys | 380 | | 385 | | 390 |
| Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gln | 395 | | 400 | | 405 |
| Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gly | 410 | | 415 | | 420 |
| Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp | 425 | | 430 | | 435 |
| Phe Pro Lys Pro Gln Ile Thr Val Gln Pro Glu Thr Gln Ser Ala | | | | | |

| | | |
|-------------------------------------|-------------------------|-----|
| 440 | 445 | 450 |
| Ile Lys Gly Ser Asn Leu Ser Phe Ile | Cys Ser Ala Ala Ser Ser | |
| 455 | 460 | 465 |
| Ser Asp Ser Pro Met Thr Phe Ala Trp | Lys Lys Asp Asn Glu Leu | |
| 470 | 475 | 480 |
| Leu His Asp Ala Glu Met Glu Asn Tyr | Ala His Leu Arg Ala Gln | |
| 485 | 490 | 495 |
| Gly Gly Glu Val Met Glu Tyr Thr Thr | Ile Leu Arg Leu Arg Glu | |
| 500 | 505 | 510 |
| Val Glu Phe Ala Ser Glu Gly Lys Tyr | Gln Cys Val Ile Ser Asn | |
| 515 | 520 | 525 |
| His Phe Gly Ser Ser Tyr Ser Val Lys | Ala Lys Leu Thr Val Asn | |
| 530 | 535 | 540 |
| Met Leu Pro Ser Phe Thr Lys Thr Pro | Met Asp Leu Thr Ile Arg | |
| 545 | 550 | 555 |
| Ala Gly Ala Met Ala Arg Leu Glu Cys | Ala Ala Val Gly His Pro | |
| 560 | 565 | 570 |
| Ala Pro Gln Ile Ala Trp Gln Lys Asp | Gly Gly Thr Asp Phe Pro | |
| 575 | 580 | 585 |
| Ala Ala Arg Glu Arg Arg Met His Val | Met Pro Glu Asp Asp Val | |
| 590 | 595 | 600 |
| Phe Phe Ile Val Asp Val Lys Ile Glu | Asp Ile Gly Val Tyr Ser | |
| 605 | 610 | 615 |
| Cys Thr Ala Gln Asn Ser Ala Gly Ser | Ile Ser Ala Asn Ala Thr | |
| 620 | 625 | 630 |
| Leu Thr Val Leu Glu Thr Pro Ser Phe | Leu Arg Pro Leu Leu Asp | |
| 635 | 640 | 645 |
| Arg Thr Val Thr Lys Gly Glu Thr Ala | Val Leu Gln Cys Ile Ala | |
| 650 | 655 | 660 |
| Gly Gly Ser Pro Pro Pro Lys Leu Asn | Trp Thr Lys Asp Asp Ser | |
| 665 | 670 | 675 |
| Pro Leu Val Val Thr Glu Arg His Phe | Phe Ala Ala Gly Asn Gln | |
| 680 | 685 | 690 |
| Leu Leu Ile Ile Val Asp Ser Asp Val | Ser Asp Ala Gly Lys Tyr | |
| 695 | 700 | 705 |
| Thr Cys Glu Met Ser Asn Thr Leu Gly | Thr Glu Arg Gly Asn Val | |
| 710 | 715 | 720 |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|-----|-----|-----|-----|------|--|
| Arg | Leu | Ser | Val | Ile | Pro | Thr | Pro | Thr | Cys | Asp | Ser | Pro | Gln | Met | |
| | | | | 725 | | | | | 730 | | | | | 735 | |
| Thr | Ala | Pro | Ser | Leu | Asp | Asp | Asp | Gly | Trp | Ala | Thr | Val | Gly | Val | |
| | | | | 740 | | | | | 745 | | | | | 750 | |
| Val | Ile | Ile | Ala | Val | Val | Cys | Cys | Val | Val | Gly | Thr | Ser | Leu | Val | |
| | | | | 755 | | | | | 760 | | | | | 765 | |
| Trp | Val | Val | Ile | Ile | Tyr | His | Thr | Arg | Arg | Arg | Asn | Glu | Asp | Cys | |
| | | | | 770 | | | | | 775 | | | | | 780 | |
| Ser | Ile | Thr | Asn | Thr | Asp | Glu | Thr | Asn | Leu | Pro | Ala | Asp | Ile | Pro | |
| | | | | 785 | | | | | 790 | | | | | 795 | |
| Ser | Tyr | Leu | Ser | Ser | Gln | Gly | Thr | Leu | Ala | Asp | Arg | Gln | Asp | Gly | |
| | | | | 800 | | | | | 805 | | | | | 810 | |
| Tyr | Val | Ser | Ser | Glu | Ser | Gly | Ser | His | His | Gln | Phe | Val | Thr | Ser | |
| | | | | 815 | | | | | 820 | | | | | 825 | |
| Ser | Gly | Ala | Gly | Phe | Phe | Leu | Pro | Gln | His | Asp | Ser | Ser | Gly | Thr | |
| | | | | 830 | | | | | 835 | | | | | 840 | |
| Cys | His | Ile | Asp | Asn | Ser | Ser | Glu | Ala | Asp | Val | Glu | Ala | Ala | Thr | |
| | | | | 845 | | | | | 850 | | | | | 855 | |
| Asp | Leu | Phe | Leu | Cys | Pro | Phe | Leu | Gly | Ser | Thr | Gly | Pro | Met | Tyr | |
| | | | | 860 | | | | | 865 | | | | | 870 | |
| Leu | Lys | Gly | Asn | Val | Tyr | Gly | Ser | Asp | Pro | Phe | Glu | Thr | Tyr | His | |
| | | | | 875 | | | | | 880 | | | | | 885 | |
| Thr | Gly | Cys | Ser | Pro | Asp | Pro | Arg | Thr | Val | Leu | Met | Asp | His | Tyr | |
| | | | | 890 | | | | | 895 | | | | | 900 | |
| Glu | Pro | Ser | Tyr | Ile | Lys | Lys | Lys | Glu | Cys | Tyr | Pro | Cys | Ser | His | |
| | | | | 905 | | | | | 910 | | | | | 915 | |
| Pro | Ser | Glu | Glu | Ser | Cys | Glu | Arg | Ser | Phe | Ser | Asn | Ile | Ser | Trp | |
| | | | | 920 | | | | | 925 | | | | | 930 | |
| Pro | Ser | His | Val | Arg | Lys | Leu | Leu | Asn | Thr | Ser | Tyr | Ser | His | Asn | |
| | | | | 935 | | | | | 940 | | | | | 945 | |
| Glu | Gly | Pro | Gly | Met | Lys | Asn | Leu | Cys | Leu | Asn | Lys | Ser | Ser | Leu | |
| | | | | 950 | | | | | 955 | | | | | 960 | |
| Asp | Phe | Ser | Ala | Asn | Pro | Glu | Pro | Ala | Ser | Val | Ala | Ser | Ser | Asn | |
| | | | | 965 | | | | | 970 | | | | | 975 | |
| Ser | Phe | Met | Gly | Thr | Phe | Gly | Lys | Ala | Leu | Arg | Arg | Pro | His | Leu | |
| | | | | 980 | | | | | 985 | | | | | 990 | |
| Asp | Ala | Tyr | Ser | Ser | Phe | Gly | Gln | Pro | Ser | Asp | Cys | Gln | Pro | Arg | |
| | | | | 995 | | | | | 1000 | | | | | 1005 | |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|------|-----|-----|-----|-----|------|-----|-----|-----|-----|------|
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| His | Ile | Cys | Thr | Phe | Lys | Gln | Thr | Leu | Glu | Asn | Tyr | Arg | Thr | Pro |
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<213> Homo Sapien

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 50 55 60
 Ile Cys Val Arg Lys Asn Leu Arg Glu Val Pro Asp Gly Ile Ser
 65 70 75
 Thr Asn Thr Arg Leu Leu Asn Leu His Glu Asn Gln Ile Gln Ile
 80 85 90
 Ile Lys Val Asn Ser Phe Lys His Leu Arg His Leu Glu Ile Leu
 95 100 105
 Gln Leu Ser Arg Asn His Ile Arg Thr Ile Glu Ile Gly Ala Phe
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| Tyr Ala Phe Asn | Arg Ile Pro Ser Leu | Arg Arg Leu Asp Leu Gly | | | |
| | 170 | | 175 | | 180 |
| Glu Leu Lys Arg | Leu Ser Tyr Ile Ser | Glu Gly Ala Phe Glu Gly | | | |
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| Leu Ser Asn Leu | Arg Tyr Leu Asn Leu | Ala Met Cys Asn Leu Arg | | | |
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| Glu Ile Pro Asn | Leu Thr Pro Leu Ile | Lys Leu Asp Glu Leu Asp | | | |
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| Leu Ser Gly Asn | His Leu Ser Ala Ile | Arg Pro Gly Ser Phe Gln | | | |
| | 230 | | 235 | | 240 |
| Gly Leu Met His | Leu Gln Lys Leu Trp | Met Ile Gln Ser Gln Ile | | | |
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| Gln Val Ile Glu | Arg Asn Ala Phe Asp | Asn Leu Gln Ser Leu Val | | | |
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| Glu Ile Asn Leu | Ala His Asn Asn Leu | Thr Leu Leu Pro His Asp | | | |
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| Leu Phe Thr Pro | Leu His His Leu Glu | Arg Ile His Leu His His | | | |
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| | 335 | | 340 | | 345 |
| Gln Asn Tyr Phe | Thr Cys Tyr Ala Pro | Val Ile Val Glu Pro Pro | | | |
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| Arg Ala Ser Thr | Ser Leu Thr Ser Val | Ser Trp Ile Thr Pro Asn | | | |
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| Gly Thr Val Met | Thr His Gly Ala Tyr | Lys Val Arg Ile Ala Val | | | |
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| | 410 | | 415 | | 420 |
| Thr Gly Met Tyr | Thr Cys Met Val Ser | Asn Ser Val Gly Asn Thr | | | |

| | 425 | | 430 | | 435 |
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| Thr Ala Ser Ala | Thr Leu Asn Val Thr | Ala Ala Thr Thr Thr | Pro | | |
| | 440 | 445 | 450 | | |
| Phe Ser Tyr Phe | Ser Thr Val Thr Val | Glu Thr Met Glu Pro | Ser | | |
| | 455 | 460 | 465 | | |
| Gln Asp Glu Ala | Arg Thr Thr Asp Asn | Asn Val Gly Pro Thr | Pro | | |
| | 470 | 475 | 480 | | |
| Val Val Asp Trp | Glu Thr Thr Asn Val | Thr Thr Ser Leu Thr | Pro | | |
| | 485 | 490 | 495 | | |
| Gln Ser Thr Arg | Ser Thr Glu Lys Thr | Phe Thr Ile Pro Val | Thr | | |
| | 500 | 505 | 510 | | |
| Asp Ile Asn Ser | Gly Ile Pro Gly Ile | Asp Glu Val Met Lys | Thr | | |
| | 515 | 520 | 525 | | |
| Thr Lys Ile Ile | Ile Gly Cys Phe Val | Ala Ile Thr Leu Met | Ala | | |
| | 530 | 535 | 540 | | |
| Ala Val Met Leu | Val Ile Phe Tyr Lys | Met Arg Lys Gln His | His | | |
| | 545 | 550 | 555 | | |
| Arg Gln Asn His | His Ala Pro Thr Arg | Thr Val Glu Ile Ile | Asn | | |
| | 560 | 565 | 570 | | |
| Val Asp Asp Glu | Ile Thr Gly Asp Thr | Pro Met Glu Ser His | Leu | | |
| | 575 | 580 | 585 | | |
| Pro Met Pro Ala | Ile Glu His Glu His | Leu Asn His Tyr Asn | Ser | | |
| | 590 | 595 | 600 | | |
| Tyr Lys Ser Pro | Phe Asn His Thr Thr | Thr Val Asn Thr Ile | Asn | | |
| | 605 | 610 | 615 | | |
| Ser Ile His Ser | Ser Val His Glu Pro | Leu Leu Ile Arg Met | Asn | | |
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<211> 4053

<212> DNA

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| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Cys | Pro | Thr | Thr | Cys | Arg | Cys | Leu | Gly | Asp | Leu | Leu | Asp | Cys |
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| Ser | Arg | Lys | Arg | Leu | Ala | Arg | Leu | Pro | Glu | Pro | Leu | Pro | Ser | Trp |
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| Val | Ala | Arg | Leu | Asp | Leu | Ser | His | Asn | Arg | Leu | Ser | Phe | Ile | Lys |
| 80 | | | | | 85 | | | | | 90 | | | | |
| Ala | Ser | Ser | Met | Ser | His | Leu | Gln | Ser | Leu | Arg | Glu | Val | Lys | Leu |
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| Asn | Asn | Asn | Glu | Leu | Glu | Thr | Ile | Pro | Asn | Leu | Gly | Pro | Val | Ser |
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| Ala | Asn | Ile | Thr | Leu | Leu | Ser | Leu | Ala | Gly | Asn | Arg | Ile | Val | Glu |
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| Ile | Leu | Pro | Glu | His | Leu | Lys | Glu | Phe | Gln | Ser | Leu | Glu | Thr | Leu |
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| Asp | Leu | Ser | Ser | Asn | Asn | Ile | Ser | Glu | Leu | Gln | Thr | Ala | Phe | Pro |
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| Ala | Leu | Gln | Leu | Lys | Tyr | Leu | Tyr | Leu | Asn | Ser | Asn | Arg | Val | Thr |
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| Ser | Met | Glu | Pro | Gly | Tyr | Phe | Asp | Asn | Leu | Ala | Asn | Thr | Leu | Leu |
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| Val | Leu | Lys | Leu | Asn | Arg | Asn | Arg | Ile | Ser | Ala | Ile | Pro | Pro | Lys |
| 200 | | | | | 205 | | | | | 210 | | | | |
| Met | Phe | Lys | Leu | Pro | Gln | Leu | Gln | His | Leu | Glu | Leu | Asn | Arg | Asn |
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| Lys | Ile | Lys | Asn | Val | Asp | Gly | Leu | Thr | Phe | Gln | Gly | Leu | Gly | Ala |
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| Leu | Lys | Ser | Leu | Lys | Met | Gln | Arg | Asn | Gly | Val | Thr | Lys | Leu | Met |
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| Asp | Gly | Ala | Phe | Trp | Gly | Leu | Ser | Asn | Met | Glu | Ile | Leu | Gln | Leu |
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| Asp | His | Asn | Asn | Leu | Thr | Glu | Ile | Thr | Lys | Gly | Trp | Leu | Tyr | Gly |
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| Leu | Leu | Met | Leu | Gln | Glu | Leu | His | Leu | Ser | Gln | Asn | Ala | Ile | Asn |
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| Arg | Ile | Ser | Pro | Asp | Ala | Trp | Glu | Phe | Cys | Gln | Lys | Leu | Ser | Glu |
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| Leu | Asp | Leu | Thr | Phe | Asn | His | Leu | Ser | Arg | Leu | Asp | Asp | Ser | Ser |

| | | | | | |
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| Arg Val Ser Tyr Ile Ala Asp Cys Ala | 350 | Phe Arg Gly Leu Ser Ser | 355 | | 360 |
| Leu Lys Thr Leu Asp Leu Lys Asn Asn | 365 | Glu Ile Ser Trp Thr Ile | 370 | | 375 |
| Glu Asp Met Asn Gly Ala Phe Ser Gly | 380 | Leu Asp Lys Leu Arg Arg | 385 | | 390 |
| Leu Ile Leu Gln Gly Asn Arg Ile Arg | 395 | Ser Ile Thr Lys Lys Ala | 400 | | 405 |
| Phe Thr Gly Leu Asp Ala Leu Glu His | 410 | Leu Asp Leu Ser Asp Asn | 415 | | 420 |
| Ala Ile Met Ser Leu Gln Gly Asn Ala | 425 | Phe Ser Gln Met Lys Lys | 430 | | 435 |
| Leu Gln Gln Leu His Leu Asn Thr Ser | 440 | Ser Leu Leu Cys Asp Cys | 445 | | 450 |
| Gln Leu Lys Trp Leu Pro Gln Trp Val | 455 | Ala Glu Asn Asn Phe Gln | 460 | | 465 |
| Ser Phe Val Asn Ala Ser Cys Ala His | 470 | Pro Gln Leu Leu Lys Gly | 475 | | 480 |
| Arg Ser Ile Phe Ala Val Ser Pro Asp | 485 | Gly Phe Val Cys Asp Asp | 490 | | 495 |
| Phe Pro Lys Pro Gln Ile Thr Val Gln | 500 | Pro Glu Thr Gln Ser Ala | 505 | | 510 |
| Ile Lys Gly Ser Asn Leu Ser Phe Ile | 515 | Cys Ser Ala Ala Ser Ser | 520 | | 525 |
| Ser Asp Ser Pro Met Thr Phe Ala Trp | 530 | Lys Lys Asp Asn Glu Leu | 535 | | 540 |
| Leu His Asp Ala Glu Met Glu Asn Tyr | 545 | Ala His Leu Arg Ala Gln | 550 | | 555 |
| Gly Gly Glu Val Met Glu Tyr Thr Thr | 560 | Ile Leu Arg Leu Arg Glu | 565 | | 570 |
| Val Glu Phe Ala Ser Glu Gly Lys Tyr | 575 | Gln Cys Val Ile Ser Asn | 580 | | 585 |
| His Phe Gly Ser Ser Tyr Ser Val Lys | 590 | Ala Lys Leu Thr Val Asn | 595 | | 600 |
| Met Leu Pro Ser Phe Thr Lys Thr Pro | | Met Asp Leu Thr Ile Arg | | | |

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| | 605 | | 610 | | 615 |
| Ala Gly Ala Met | Ala Arg Leu Glu Cys | Ala Ala Val Gly His | Pro | | |
| | 620 | | 625 | | 630 |
| Ala Pro Gln Ile | Ala Trp Gln Lys Asp | Gly Gly Thr Asp Phe | Pro | | |
| | 635 | | 640 | | 645 |
| Ala Ala Arg Glu | Arg Arg Met His Val | Met Pro Glu Asp Asp | Val | | |
| | 650 | | 655 | | 660 |
| Phe Phe Ile Val | Asp Val Lys Ile Glu | Asp Ile Gly Val Tyr | Ser | | |
| | 665 | | 670 | | 675 |
| Cys Thr Ala Gln | Asn Ser Ala Gly Ser | Ile Ser Ala Asn Ala | Thr | | |
| | 680 | | 685 | | 690 |
| Leu Thr Val Leu | Glu Thr Pro Ser Phe | Leu Arg Pro Leu Leu | Asp | | |
| | 695 | | 700 | | 705 |
| Arg Thr Val Thr | Lys Gly Glu Thr Ala | Val Leu Gln Cys Ile | Ala | | |
| | 710 | | 715 | | 720 |
| Gly Gly Ser Pro | Pro Pro Lys Leu Asn | Trp Thr Lys Asp Asp | Ser | | |
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| Pro Leu Val Val | Thr Glu Arg His Phe | Phe Ala Ala Gly Asn | Gln | | |
| | 740 | | 745 | | 750 |
| Leu Leu Ile Ile | Val Asp Ser Asp Val | Ser Asp Ala Gly Lys | Tyr | | |
| | 755 | | 760 | | 765 |
| Thr Cys Glu Met | Ser Asn Thr Leu Gly | Thr Glu Arg Gly Asn | Val | | |
| | 770 | | 775 | | 780 |
| Arg Leu Ser Val | Ile Pro Thr Pro Thr | Cys Asp Ser Pro Gln | Met | | |
| | 785 | | 790 | | 795 |
| Thr Ala Pro Ser | Leu Asp Asp Asp Gly | Trp Ala Thr Val Gly | Val | | |
| | 800 | | 805 | | 810 |
| Val Ile Ile Ala | Val Val Cys Cys Val | Val Gly Thr Ser Leu | Val | | |
| | 815 | | 820 | | 825 |
| Trp Val Val Ile | Ile Tyr His Thr Arg | Arg Arg Asn Glu Asp | Cys | | |
| | 830 | | 835 | | 840 |
| Ser Ile Thr Asn | Thr Asp Glu Thr Asn | Leu Pro Ala Asp Ile | Pro | | |
| | 845 | | 850 | | 855 |
| Ser Tyr Leu Ser | Ser Gln Gly Thr Leu | Ala Asp Arg Gln Asp | Gly | | |
| | 860 | | 865 | | 870 |
| Tyr Val Ser Ser | Glu Ser Gly Ser His | His Gln Phe Val Thr | Ser | | |
| | 875 | | 880 | | 885 |
| Ser Gly Ala Gly | Phe Phe Leu Pro Gln | His Asp Ser Ser Gly | Thr | | |

| 890 | | | | | | | | | | 895 | | | | | 900 | | | | |
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| Cys | His | Ile | Asp | Asn | Ser | Ser | Glu | Ala | Asp | Val | Glu | Ala | Ala | Thr | | | | | |
| | | | | 905 | | | | | 910 | | | | | 915 | | | | | |
| Asp | Leu | Phe | Leu | Cys | Pro | Phe | Leu | Gly | Ser | Thr | Gly | Pro | Met | Tyr | | | | | |
| | | | | 920 | | | | | 925 | | | | | 930 | | | | | |
| Leu | Lys | Gly | Asn | Val | Tyr | Gly | Ser | Asp | Pro | Phe | Glu | Thr | Tyr | His | | | | | |
| | | | | 935 | | | | | 940 | | | | | 945 | | | | | |
| Thr | Gly | Cys | Ser | Pro | Asp | Pro | Arg | Thr | Val | Leu | Met | Asp | His | Tyr | | | | | |
| | | | | 950 | | | | | 955 | | | | | 960 | | | | | |
| Glu | Pro | Ser | Tyr | Ile | Lys | Lys | Lys | Glu | Cys | Tyr | Pro | Cys | Ser | His | | | | | |
| | | | | 965 | | | | | 970 | | | | | 975 | | | | | |
| Pro | Ser | Glu | Glu | Ser | Cys | Glu | Arg | Ser | Phe | Ser | Asn | Ile | Ser | Trp | | | | | |
| | | | | 980 | | | | | 985 | | | | | 990 | | | | | |
| Pro | Ser | His | Val | Arg | Lys | Leu | Leu | Asn | Thr | Ser | Tyr | Ser | His | Asn | | | | | |
| | | | | 995 | | | | | 1000 | | | | | 1005 | | | | | |
| Glu | Gly | Pro | Gly | Met | Lys | Asn | Leu | Cys | Leu | Asn | Lys | Ser | Ser | Leu | | | | | |
| | | | | 1010 | | | | | 1015 | | | | | 1020 | | | | | |
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| Ser | Phe | Met | Gly | Thr | Phe | Gly | Lys | Ala | Leu | Arg | Arg | Pro | His | Leu | | | | | |
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| | | | | 1070 | | | | | 1075 | | | | | 1080 | | | | | |
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| | | | | 1085 | | | | | 1090 | | | | | 1095 | | | | | |
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| Met | Asp | Phe | Leu | Leu | Ala | Leu | Val | Leu | Val | Ser | Ser | Leu | Tyr | Leu | 1 | 5 | 10 | 15 |
| Gln | Ala | Ala | Ala | Glu | Phe | Asp | Gly | Arg | Trp | Pro | Arg | Gln | Ile | Val | 20 | 25 | 30 | |
| Ser | Ser | Ile | Gly | Leu | Cys | Arg | Tyr | Gly | Gly | Arg | Ile | Asp | Cys | Cys | 35 | 40 | 45 | |
| Trp | Gly | Trp | Ala | Arg | Gln | Ser | Trp | Gly | Gln | Cys | Gln | Pro | Val | Cys | 50 | 55 | 60 | |
| Gln | Pro | Arg | Cys | Lys | His | Gly | Glu | Cys | Ile | Gly | Pro | Asn | Lys | Cys | 65 | 70 | 75 | |
| Lys | Cys | His | Pro | Gly | Tyr | Ala | Gly | Lys | Thr | Cys | Asn | Gln | Asp | Leu | 80 | 85 | 90 | |
| Asn | Glu | Cys | Gly | Leu | Lys | Pro | Arg | Pro | Cys | Lys | His | Arg | Cys | Met | 95 | 100 | 105 | |
| Asn | Thr | Tyr | Gly | Ser | Tyr | Lys | Cys | Tyr | Cys | Leu | Asn | Gly | Tyr | Met | 110 | 115 | 120 | |
| Leu | Met | Pro | Asp | Gly | Ser | Cys | Ser | Ser | Ala | Leu | Thr | Cys | Ser | Met | 125 | 130 | 135 | |
| Ala | Asn | Cys | Gln | Tyr | Gly | Cys | Asp | Val | Val | Lys | Gly | Gln | Ile | Arg | 140 | 145 | 150 | |
| Cys | Gln | Cys | Pro | Ser | Pro | Gly | Leu | His | Leu | Ala | Pro | Asp | Gly | Arg | 155 | 160 | 165 | |
| Thr | Cys | Val | Asp | Val | Asp | Glu | Cys | Ala | Thr | Gly | Arg | Ala | Ser | Cys | 170 | 175 | 180 | |
| Pro | Arg | Phe | Arg | Gln | Cys | Val | Asn | Thr | Phe | Gly | Ser | Tyr | Ile | Cys | 185 | 190 | 195 | |
| Lys | Cys | His | Lys | Gly | Phe | Asp | Leu | Met | Tyr | Ile | Gly | Gly | Lys | Tyr | 200 | 205 | 210 | |
| Gln | Cys | His | Asp | Ile | Asp | Glu | Cys | Ser | Leu | Gly | Gln | Tyr | Gln | Cys | 215 | 220 | 225 | |
| Ser | Ser | Phe | Ala | Arg | Cys | Tyr | Asn | Val | Arg | Gly | Ser | Tyr | Lys | Cys | 230 | 235 | 240 | |
| Lys | Cys | Lys | Glu | Gly | Tyr | Gln | Gly | Asp | Gly | Leu | Thr | Cys | Val | Tyr | | | | |

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| Lys | Gly | Asn | Gly | Thr | Ile | Leu | Lys | Gly | Asp | Thr | Gly | Asn | Asn | Asn |
| | | | | 275 | | | | . | 280 | | | | | 285 |
| Trp | Ile | Pro | Asp | Val | Gly | Ser | Thr | Trp | Trp | Pro | Pro | Lys | Thr | Pro |
| | | | | 290 | | | | | 295 | | | | | 300 |
| Tyr | Ile | Pro | Pro | Ile | Ile | Thr | Asn | Arg | Pro | Thr | Ser | Lys | Pro | Thr |
| | | | | 305 | | | | | 310 | | | | | 315 |
| Thr | Arg | Pro | Thr | Pro | Lys | Pro | Thr | Pro | Ile | Pro | Thr | Pro | Pro | Pro |
| | | | | 320 | | | | | 325 | | | | | 330 |
| Pro | Pro | Pro | Leu | Pro | Thr | Glu | Leu | Arg | Thr | Pro | Leu | Pro | Pro | Thr |
| | | | | 335 | | | | | 340 | | | | | 345 |
| Thr | Pro | Glu | Arg | Pro | Thr | Thr | Gly | Leu | Thr | Thr | Ile | Ala | Pro | Ala |
| | | | | 350 | | | | | 355 | | | | | 360 |
| Ala | Ser | Thr | Pro | Pro | Gly | Gly | Ile | Thr | Val | Asp | Asn | Arg | Val | Gln |
| | | | | 365 | | | | | 370 | | | | | 375 |
| Thr | Asp | Pro | Gln | Lys | Pro | Arg | Gly | Asp | Val | Phe | Ser | Val | Leu | Val |
| | | | | 380 | | | | | 385 | | | | | 390 |
| His | Ser | Cys | Asn | Phe | Asp | His | Gly | Leu | Cys | Gly | Trp | Ile | Arg | Glu |
| | | | | 395 | | | | | 400 | | | | | 405 |
| Lys | Asp | Asn | Asp | Leu | His | Trp | Glu | Pro | Ile | Arg | Asp | Pro | Ala | Gly |
| | | | | 410 | | | | | 415 | | | | | 420 |
| Gly | Gln | Tyr | Leu | Thr | Val | Ser | Ala | Ala | Lys | Ala | Pro | Gly | Gly | Lys |
| | | | | 425 | | | | | 430 | | | | | 435 |
| Ala | Ala | Arg | Leu | Val | Leu | Pro | Leu | Gly | Arg | Leu | Met | His | Ser | Gly |
| | | | | 440 | | | | | 445 | | | | | 450 |
| Asp | Leu | Cys | Leu | Ser | Phe | Arg | His | Lys | Val | Thr | Gly | Leu | His | Ser |
| | | | | 455 | | | | | 460 | | | | | 465 |
| Gly | Thr | Leu | Gln | Val | Phe | Val | Arg | Lys | His | Gly | Ala | His | Gly | Ala |
| | | | | 470 | | | | | 475 | | | | | 480 |
| Ala | Leu | Trp | Gly | Arg | Asn | Gly | Gly | His | Gly | Trp | Arg | Gln | Thr | Gln |
| | | | | 485 | | | | | 490 | | | | | 495 |
| Ile | Thr | Leu | Arg | Gly | Ala | Asp | Ile | Lys | Ser | Glu | Ser | Gln | Arg | |
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<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 316

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<210> 317

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 317

ttgcacttgt aggaccacg tacg 24

<210> 318

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 318

ctgatgggag gacctgtgta gatgttgatg aatgtgctac aggaagagcc 50

<210> 319

<211> 2110

<212> DNA

<213> Homo Sapien

<400> 319

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tttagattgt gaaatgtggc tcaaggtctt cacaactttc ctttcctttg 100

caacaggtgc ttgctcgggg ctgaaggtga cagtgccatc acacactgtc 150

catggcgtca gaggtcaggc cctctaccta cccgtccact atggcttcca 200

cactccagca tcagacatcc agatcatatg gctatttgag agaccccaca 250

caatgccc aaatacttactg ggctctgtga ataagtctgt gggttcctgac 300

ttggaatacc aacacaagtt caccatgatg ccacccaatg catctctgct 350

tatcaaccca ctgcagttcc ctgatgaagg caattacatc gtgaaggtca 400

acattcaggg aaatggaact ctatctgcca gtcagaagat acaagtcacg 450

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aaaaaaaaa 2110

<210> 320
<211> 450
<212> PRT
<213> Homo Sapien

<400> 320

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Met | Trp | Leu | Lys | Val | Phe | Thr | Thr | Phe | Leu | Ser | Phe | Ala | Thr | Gly | |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Ala | Cys | Ser | Gly | Leu | Lys | Val | Thr | Val | Pro | Ser | His | Thr | Val | His | |
| | | | | 20 | | | | | 25 | | | | | 30 | |
| Gly | Val | Arg | Gly | Gln | Ala | Leu | Tyr | Leu | Pro | Val | His | Tyr | Gly | Phe | |
| | | | | 35 | | | | | 40 | | | | | 45 | |
| His | Thr | Pro | Ala | Ser | Asp | Ile | Gln | Ile | Ile | Trp | Leu | Phe | Glu | Arg | |
| | | | | 50 | | | | | 55 | | | | | 60 | |
| Pro | His | Thr | Met | Pro | Lys | Tyr | Leu | Leu | Gly | Ser | Val | Asn | Lys | Ser | |
| | | | | 65 | | | | | 70 | | | | | 75 | |
| Val | Val | Pro | Asp | Leu | Glu | Tyr | Gln | His | Lys | Phe | Thr | Met | Met | Pro | |
| | | | | 80 | | | | | 85 | | | | | 90 | |
| Pro | Asn | Ala | Ser | Leu | Leu | Ile | Asn | Pro | Leu | Gln | Phe | Pro | Asp | Glu | |
| | | | | 95 | | | | | 100 | | | | | 105 | |
| Gly | Asn | Tyr | Ile | Val | Lys | Val | Asn | Ile | Gln | Gly | Asn | Gly | Thr | Leu | |
| | | | | 110 | | | | | 115 | | | | | 120 | |
| Ser | Ala | Ser | Gln | Lys | Ile | Gln | Val | Thr | Val | Asp | Asp | Pro | Val | Thr | |
| | | | | 125 | | | | | 130 | | | | | 135 | |
| Lys | Pro | Val | Val | Gln | Ile | His | Pro | Pro | Ser | Gly | Ala | Val | Glu | Tyr | |
| | | | | 140 | | | | | 145 | | | | | 150 | |
| Val | Gly | Asn | Met | Thr | Leu | Thr | Cys | His | Val | Glu | Gly | Gly | Thr | Arg | |
| | | | | 155 | | | | | 160 | | | | | 165 | |
| Leu | Ala | Tyr | Gln | Trp | Leu | Lys | Asn | Gly | Arg | Pro | Val | His | Thr | Ser | |
| | | | | 170 | | | | | 175 | | | | | 180 | |
| Ser | Thr | Tyr | Ser | Phe | Ser | Pro | Gln | Asn | Asn | Thr | Leu | His | Ile | Ala | |
| | | | | 185 | | | | | 190 | | | | | 195 | |
| Pro | Val | Thr | Lys | Glu | Asp | Ile | Gly | Asn | Tyr | Ser | Cys | Leu | Val | Arg | |
| | | | | 200 | | | | | 205 | | | | | 210 | |
| Asn | Pro | Val | Ser | Glu | Met | Glu | Ser | Asp | Ile | Ile | Met | Pro | Ile | Ile | |

| 215 | | | | | 220 | | | | | 225 | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Tyr | Tyr | Gly | Pro | Tyr | Gly | Leu | Gln | Val | Asn | Ser | Asp | Lys | Gly | Leu |
| | | | | 230 | | | | | 235 | | | | | 240 |
| Lys | Val | Gly | Glu | Val | Phe | Thr | Val | Asp | Leu | Gly | Glu | Ala | Ile | Leu |
| | | | | 245 | | | | | 250 | | | | | 255 |
| Phe | Asp | Cys | Ser | Ala | Asp | Ser | His | Pro | Pro | Asn | Thr | Tyr | Ser | Trp |
| | | | | 260 | | | | | 265 | | | | | 270 |
| Ile | Arg | Arg | Thr | Asp | Asn | Thr | Thr | Tyr | Ile | Ile | Lys | His | Gly | Pro |
| | | | | 275 | | | | | 280 | | | | | 285 |
| Arg | Leu | Glu | Val | Ala | Ser | Glu | Lys | Val | Ala | Gln | Lys | Thr | Met | Asp |
| | | | | 290 | | | | | 295 | | | | | 300 |
| Tyr | Val | Cys | Cys | Ala | Tyr | Asn | Asn | Ile | Thr | Gly | Arg | Gln | Asp | Glu |
| | | | | 305 | | | | | 310 | | | | | 315 |
| Thr | His | Phe | Thr | Val | Ile | Ile | Thr | Ser | Val | Gly | Leu | Glu | Lys | Leu |
| | | | | 320 | | | | | 325 | | | | | 330 |
| Ala | Gln | Lys | Gly | Lys | Ser | Leu | Ser | Pro | Leu | Ala | Ser | Ile | Thr | Gly |
| | | | | 335 | | | | | 340 | | | | | 345 |
| Ile | Ser | Leu | Phe | Leu | Ile | Ile | Ser | Met | Cys | Leu | Leu | Phe | Leu | Trp |
| | | | | 350 | | | | | 355 | | | | | 360 |
| Lys | Lys | Tyr | Gln | Pro | Tyr | Lys | Val | Ile | Lys | Gln | Lys | Leu | Glu | Gly |
| | | | | 365 | | | | | 370 | | | | | 375 |
| Arg | Pro | Glu | Thr | Glu | Tyr | Arg | Lys | Ala | Gln | Thr | Phe | Ser | Gly | His |
| | | | | 380 | | | | | 385 | | | | | 390 |
| Glu | Asp | Ala | Leu | Asp | Asp | Phe | Gly | Ile | Tyr | Glu | Phe | Val | Ala | Phe |
| | | | | 395 | | | | | 400 | | | | | 405 |
| Pro | Asp | Val | Ser | Gly | Val | Ser | Arg | Ile | Pro | Ser | Arg | Ser | Val | Pro |
| | | | | 410 | | | | | 415 | | | | | 420 |
| Ala | Ser | Asp | Cys | Val | Ser | Gly | Gln | Asp | Leu | His | Ser | Thr | Val | Tyr |
| | | | | 425 | | | | | 430 | | | | | 435 |
| Glu | Val | Ile | Gln | His | Ile | Pro | Ala | Gln | Gln | Gln | Asp | His | Pro | Glu |
| | | | | 440 | | | | | 445 | | | | | 450 |

<210> 321

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 321

gatacctgtca caaagccagt ggtgc 25

<210> 322

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 322

cactgacagg gttcctcacc cagg 24

<210> 323

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 323

ctccctctgg gctgtggagt atgtggggaa catgaccctg acatg 45

<210> 324

<211> 2397

<212> DNA

<213> Homo Sapien

<400> 324

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cctggtgctg ttgctttggg gtgctccctg gacgcacggg cggcggagca 100
acgttcgcgt catcacggac gagaactgga gagaactgct ggaaggagac 150
tggatgatag aattttatgc cccgtggtgc cctgcttgtc aaaatcttca 200
accggaatgg gaaagttttg ctgaatgggg agaagatctt gaggttaata 250
ttgcgaaagt agatgtcaca gacgagccag gactgagtgg acggtttatc 300
ataactgctc ttcctactat ttatcattgt aaagatggtg aatttaggcg 350
ctatcagggg ccaaggacta agaaggactt cataaacttt ataagtgata 400
aagagtggaa gagtattgag cccgtttcat catggtttgg tccaggttct 450
gttctgatga gtagtatgtc agcactcttt cagctatcta tgtggatcag 500
gacgtgccat aactacttta ttgaagacct tggattgcca gtgtggggat 550
catatactgt ttttgcttta gcaactctgt tttccggact gttattagga 600
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accacagcca taccataacc cttcaaaaaa attattatca gaatctgcac 700

aacctttgaa aaaagtggag gaggaacaag aggcggatga agaagatggt 750
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gaatgccata agacaacgct ctctgggtcc atcattggcc acagataaat 850
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 ttcttttaaag ccctctcctt tagaatttaa aatattgtac cattaaagag 2300
 tttggatgtg taacttgtaga tgccttagaa aaataticta agcacaaaat 2350
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<210> 325

<211> 280

<212> PRT

<213> Homo Sapien

<400> 325

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|
| Met | Ala | Pro | Ser | Gly | Ser | Leu | Ala | Val | Pro | Leu | Ala | Val | Leu | Val | 1 | 5 | 10 | 15 |
| Leu | Leu | Leu | Trp | Gly | Ala | Pro | Trp | Thr | His | Gly | Arg | Arg | Ser | Asn | 20 | 25 | 30 | |
| Val | Arg | Val | Ile | Thr | Asp | Glu | Asn | Trp | Arg | Glu | Leu | Leu | Glu | Gly | 35 | 40 | 45 | |
| Asp | Trp | Met | Ile | Glu | Phe | Tyr | Ala | Pro | Trp | Cys | Pro | Ala | Cys | Gln | 50 | 55 | 60 | |
| Asn | Leu | Gln | Pro | Glu | Trp | Glu | Ser | Phe | Ala | Glu | Trp | Gly | Glu | Asp | 65 | 70 | 75 | |
| Leu | Glu | Val | Asn | Ile | Ala | Lys | Val | Asp | Val | Thr | Glu | Gln | Pro | Gly | 80 | 85 | 90 | |
| Leu | Ser | Gly | Arg | Phe | Ile | Ile | Thr | Ala | Leu | Pro | Thr | Ile | Tyr | His | 95 | 100 | 105 | |
| Cys | Lys | Asp | Gly | Glu | Phe | Arg | Arg | Tyr | Gln | Gly | Pro | Arg | Thr | Lys | 110 | 115 | 120 | |
| Lys | Asp | Phe | Ile | Asn | Phe | Ile | Ser | Asp | Lys | Glu | Trp | Lys | Ser | Ile | 125 | 130 | 135 | |
| Glu | Pro | Val | Ser | Ser | Trp | Phe | Gly | Pro | Gly | Ser | Val | Leu | Met | Ser | 140 | 145 | 150 | |
| Ser | Met | Ser | Ala | Leu | Phe | Gln | Leu | Ser | Met | Trp | Ile | Arg | Thr | Cys | 155 | 160 | 165 | |
| His | Asn | Tyr | Phe | Ile | Glu | Asp | Leu | Gly | Leu | Pro | Val | Trp | Gly | Ser | 170 | 175 | 180 | |
| Tyr | Thr | Val | Phe | Ala | Leu | Ala | Thr | Leu | Phe | Ser | Gly | Leu | Leu | Leu | 185 | 190 | 195 | |

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Leu | Cys | Met | Ile | Phe | Val | Ala | Asp | Cys | Leu | Cys | Pro | Ser | Lys | 200 | 205 | 210 |
| Arg | Arg | Arg | Pro | Gln | Pro | Tyr | Pro | Tyr | Pro | Ser | Lys | Lys | Leu | Leu | 215 | 220 | 225 |
| Ser | Glu | Ser | Ala | Gln | Pro | Leu | Lys | Lys | Val | Glu | Glu | Glu | Gln | Glu | 230 | 235 | 240 |
| Ala | Asp | Glu | Glu | Asp | Val | Ser | Glu | Glu | Glu | Ala | Glu | Ser | Lys | Glu | 245 | 250 | 255 |
| Gly | Thr | Asn | Lys | Asp | Phe | Pro | Gln | Asn | Ala | Ile | Arg | Gln | Arg | Ser | 260 | 265 | 270 |
| Leu | Gly | Pro | Ser | Leu | Ala | Thr | Asp | Lys | Ser | | | | | | 275 | 280 | |

<210> 326

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 326

tgaggtgggc aagcggcgaa atg 23

<210> 327

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 327

tatgtggatc aggacgtgcc 20

<210> 328

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 328

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<210> 329

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 329

ttgaaggaca aaggcaatct gccac 25

<210> 330

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 330

ggagtcttg gc agttcccctg gcagtcctgg tgctgttgct ttggg 45

<210> 331

<211> 2168

<212> DNA

<213> Homo Sapien

<400> 331

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aacgggaccc ttctgtgtgc cagaaaccgc aagcagttgc taaccagtg 100

ggacaggcgg attggaagag cgggaaggct ctggcccaga gcagtgtgac 150

acttccctct gtgaccatga aactctgggt gtctgcattg ctgatggcct 200

ggtttggtgt cctgagctgt gtgcaggccg aattcttcac ctctattggg 250

cacatgactg acctgattta tgcaagaaaa gagctggtgc agtctctgaa 300

agagtacatc cttgtggagg aagccaagct ttccaagatt aagagctggg 350

ccaacaaaat ggaagccttg actagcaagt cagctgctga tgctgagggc 400

tacctggctc accctgtgaa tgcctacaaa ctggtgaagc ggctaaacac 450

agactggcct gcgctggagg acctgtcct gcaggactca gctgcagggt 500

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gagataggag ctgccaagc cctgatgaga cttcaggaca catacaggct 600

ggaccagggc acaatttcca gaggggaact tccaggaacc aagtaccagg 650

caatgctgag tgtggatgac tgctttggga tgggccgctc ggctacaat 700

gaaggggact attatcatc ggtgttgtgg atggagcagg tgctaaagca 750

gcttgatgcc ggggaggagg ccaccacaac caagtcacag gtgctggact 800

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<213> Homo Sapien

<400> 332

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| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Leu | Ser | Cys | Val | Gln | Ala | Glu | Phe | Phe | Thr | Ser | Ile | Gly | His | Met | |
| | | | | 20 | | | | | 25 | | | | | 30 | |
| Thr | Asp | Leu | Ile | Tyr | Ala | Glu | Lys | Glu | Leu | Val | Gln | Ser | Leu | Lys | |
| | | | | 35 | | | | | 40 | | | | | 45 | |
| Glu | Tyr | Ile | Leu | Val | Glu | Glu | Ala | Lys | Leu | Ser | Lys | Ile | Lys | Ser | |
| | | | | 50 | | | | | 55 | | | | | 60 | |
| Trp | Ala | Asn | Lys | Met | Glu | Ala | Leu | Thr | Ser | Lys | Ser | Ala | Ala | Asp | |
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| Ala | Glu | Gly | Tyr | Leu | Ala | His | Pro | Val | Asn | Ala | Tyr | Lys | Leu | Val | |
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| Lys | Arg | Leu | Asn | Thr | Asp | Trp | Pro | Ala | Leu | Glu | Asp | Leu | Val | Leu | |
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| Gln | Asp | Ser | Ala | Ala | Gly | Phe | Ile | Ala | Asn | Leu | Ser | Val | Gln | Arg | |
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| Gln | Phe | Phe | Pro | Thr | Asp | Glu | Asp | Glu | Ile | Gly | Ala | Ala | Lys | Ala | |
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| Leu | Met | Arg | Leu | Gln | Asp | Thr | Tyr | Arg | Leu | Asp | Pro | Gly | Thr | Ile | |
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| Ser | Arg | Gly | Glu | Leu | Pro | Gly | Thr | Lys | Tyr | Gln | Ala | Met | Leu | Ser | |
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| Val | Asp | Asp | Cys | Phe | Gly | Met | Gly | Arg | Ser | Ala | Tyr | Asn | Glu | Gly | |
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| Asp | Tyr | Tyr | His | Thr | Val | Leu | Trp | Met | Glu | Gln | Val | Leu | Lys | Gln | |
| | | | | 185 | | | | | 190 | | | | | 195 | |
| Leu | Asp | Ala | Gly | Glu | Glu | Ala | Thr | Thr | Thr | Lys | Ser | Gln | Val | Leu | |
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| Asp | Tyr | Leu | Ser | Tyr | Ala | Val | Phe | Gln | Leu | Gly | Asp | Leu | His | Arg | |
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| Ala | Leu | Glu | Leu | Thr | Arg | Arg | Leu | Leu | Ser | Leu | Asp | Pro | Ser | His | |
| | | | | 230 | | | | | 235 | | | | | 240 | |
| Glu | Arg | Ala | Gly | Gly | Asn | Leu | Arg | Tyr | Phe | Glu | Gln | Leu | Leu | Glu | |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Glu | Glu | Arg | Glu | Lys | Thr | Leu | Thr | Asn | Gln | Thr | Glu | Ala | Glu | Leu | |
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| | | | |
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| Ala Thr Pro Glu Gly Ile Tyr Glu Arg Pro Val Asp Tyr Leu Pro | 275 | 280 | 285 |
| Glu Arg Asp Val Tyr Glu Ser Leu Cys Arg Gly Glu Gly Val Lys | 290 | 295 | 300 |
| Leu Thr Pro Arg Arg Gln Lys Arg Leu Phe Cys Arg Tyr His His | 305 | 310 | 315 |
| Gly Asn Arg Ala Pro Gln Leu Leu Ile Ala Pro Phe Lys Glu Glu | 320 | 325 | 330 |
| Asp Glu Trp Asp Ser Pro His Ile Val Arg Tyr Tyr Asp Val Met | 335 | 340 | 345 |
| Ser Asp Glu Glu Ile Glu Arg Ile Lys Glu Ile Ala Lys Pro Lys | 350 | 355 | 360 |
| Leu Ala Arg Ala Thr Val Arg Asp Pro Lys Thr Gly Val Leu Thr | 365 | 370 | 375 |
| Val Ala Ser Tyr Arg Val Ser Lys Ser Ser Trp Leu Glu Glu Asp | 380 | 385 | 390 |
| Asp Asp Pro Val Val Ala Arg Val Asn Arg Arg Met Gln His Ile | 395 | 400 | 405 |
| Thr Gly Leu Thr Val Lys Thr Ala Glu Leu Leu Gln Val Ala Asn | 410 | 415 | 420 |
| Tyr Gly Val Gly Gly Gln Tyr Glu Pro His Phe Asp Phe Ser Arg | 425 | 430 | 435 |
| Arg Pro Phe Asp Ser Gly Leu Lys Thr Glu Gly Asn Arg Leu Ala | 440 | 445 | 450 |
| Thr Phe Leu Asn Tyr Met Ser Asp Val Glu Ala Gly Gly Ala Thr | 455 | 460 | 465 |
| Val Phe Pro Asp Leu Gly Ala Ala Ile Trp Pro Lys Lys Gly Thr | 470 | 475 | 480 |
| Ala Val Phe Trp Tyr Asn Leu Leu Arg Ser Gly Glu Gly Asp Tyr | 485 | 490 | 495 |
| Arg Thr Arg His Ala Ala Cys Pro Val Leu Val Gly Cys Lys Trp | 500 | 505 | 510 |
| Val Ser Asn Lys Trp Phe His Glu Arg Gly Gln Glu Phe Leu Arg | 515 | 520 | 525 |
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<400> 339

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| Met | Arg | Leu | Ser | Ser | Leu | Leu | Ala | Leu | Leu | Arg | Pro | Ala | Leu | Pro | 1 | 5 | 10 | 15 |
| Leu | Ile | Leu | Gly | Leu | Ser | Leu | Gly | Cys | Ser | Leu | Ser | Leu | Leu | Arg | 20 | 25 | 30 | |
| Val | Ser | Trp | Ile | Gln | Gly | Glu | Gly | Glu | Asp | Pro | Cys | Val | Glu | Ala | 35 | 40 | 45 | |
| Val | Gly | Glu | Arg | Gly | Gly | Pro | Gln | Asn | Pro | Asp | Ser | Arg | Ala | Arg | 50 | 55 | 60 | |
| Leu | Asp | Gln | Ser | Asp | Glu | Asp | Phe | Lys | Pro | Arg | Ile | Val | Pro | Tyr | 65 | 70 | 75 | |
| Tyr | Arg | Asp | Pro | Asn | Lys | Pro | Tyr | Lys | Lys | Val | Leu | Arg | Thr | Arg | 80 | 85 | 90 | |
| Tyr | Ile | Gln | Thr | Glu | Leu | Gly | Ser | Arg | Glu | Arg | Leu | Leu | Val | Ala | 95 | 100 | 105 | |
| Val | Leu | Thr | Ser | Arg | Ala | Thr | Leu | Ser | Thr | Leu | Ala | Val | Ala | Val | 110 | 115 | 120 | |
| Asn | Arg | Thr | Val | Ala | His | His | Phe | Pro | Arg | Leu | Leu | Tyr | Phe | Thr | 125 | 130 | 135 | |
| Gly | Gln | Arg | Gly | Ala | Arg | Ala | Pro | Ala | Gly | Met | Gln | Val | Val | Ser | 140 | 145 | 150 | |
| His | Gly | Asp | Glu | Arg | Pro | Ala | Trp | Leu | Met | Ser | Glu | Thr | Leu | Arg | 155 | 160 | 165 | |
| His | Leu | His | Thr | His | Phe | Gly | Ala | Asp | Tyr | Asp | Trp | Phe | Phe | Ile | 170 | 175 | 180 | |
| Met | Gln | Asp | Asp | Thr | Tyr | Val | Gln | Ala | Pro | Arg | Leu | Ala | Ala | Leu | 185 | 190 | 195 | |
| Ala | Gly | His | Leu | Ser | Ile | Asn | Gln | Asp | Leu | Tyr | Leu | Gly | Arg | Ala | 200 | 205 | 210 | |
| Glu | Glu | Phe | Ile | Gly | Ala | Gly | Glu | Gln | Ala | Arg | Tyr | Cys | His | Gly | 215 | 220 | 225 | |
| Gly | Phe | Gly | Tyr | Leu | Leu | Ser | Arg | Ser | Leu | Leu | Leu | Arg | Leu | Arg | 230 | 235 | 240 | |
| Pro | His | Leu | Asp | Gly | Cys | Arg | Gly | Asp | Ile | Leu | Ser | Ala | Arg | Pro | 245 | 250 | 255 | |

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|-----|-----|-----|
| Asp | Glu | Trp | Leu | Gly | Arg | Cys | Leu | Ile | Asp | Ser | Leu | Gly | Val | Gly | | 260 | 265 | 270 |
| Cys | Val | Ser | Gln | His | Gln | Gly | Gln | Gln | Tyr | Arg | Ser | Phe | Glu | Leu | | 275 | 280 | 285 |
| Ala | Lys | Asn | Arg | Asp | Pro | Glu | Lys | Glu | Gly | Ser | Ser | Ala | Phe | Leu | | 290 | 295 | 300 |
| Ser | Ala | Phe | Ala | Val | His | Pro | Val | Ser | Glu | Gly | Thr | Leu | Met | Tyr | | 305 | 310 | 315 |
| Arg | Leu | His | Lys | Arg | Phe | Ser | Ala | Leu | Glu | Leu | Glu | Arg | Ala | Tyr | | 320 | 325 | 330 |
| Ser | Glu | Ile | Glu | Gln | Leu | Gln | Ala | Gln | Ile | Arg | Asn | Leu | Thr | Val | | 335 | 340 | 345 |
| Leu | Thr | Pro | Glu | Gly | Glu | Ala | Gly | Leu | Ser | Trp | Pro | Val | Gly | Leu | | 350 | 355 | 360 |
| Pro | Ala | Pro | Phe | Thr | Pro | His | Ser | Arg | Phe | Glu | Val | Leu | Gly | Trp | | 365 | 370 | 375 |
| Asp | Tyr | Phe | Thr | Glu | Gln | His | Thr | Phe | Ser | Cys | Ala | Asp | Gly | Ala | | 380 | 385 | 390 |
| Pro | Lys | Cys | Pro | Leu | Gln | Gly | Ala | Ser | Arg | Ala | Asp | Val | Gly | Asp | | 395 | 400 | 405 |
| Ala | Leu | Glu | Thr | Ala | Leu | Glu | Gln | Leu | Asn | Arg | Arg | Tyr | Gln | Pro | | 410 | 415 | 420 |
| Arg | Leu | Arg | Phe | Gln | Lys | Gln | Arg | Leu | Leu | Asn | Gly | Tyr | Arg | Arg | | 425 | 430 | 435 |
| Phe | Asp | Pro | Ala | Arg | Gly | Met | Glu | Tyr | Thr | Leu | Asp | Leu | Leu | Leu | | 440 | 445 | 450 |
| Glu | Cys | Val | Thr | Gln | Arg | Gly | His | Arg | Arg | Ala | Leu | Ala | Arg | Arg | | 455 | 460 | 465 |
| Val | Ser | Leu | Leu | Arg | Pro | Leu | Ser | Arg | Val | Glu | Ile | Leu | Pro | Met | | 470 | 475 | 480 |
| Pro | Tyr | Val | Thr | Glu | Ala | Thr | Arg | Val | Gln | Leu | Val | Leu | Pro | Leu | | 485 | 490 | 495 |
| Leu | Val | Ala | Glu | Ala | Ala | Ala | Ala | Pro | Ala | Phe | Leu | Glu | Ala | Phe | | 500 | 505 | 510 |
| Ala | Ala | Asn | Val | Leu | Glu | Pro | Arg | Glu | His | Ala | Leu | Leu | Thr | Leu | | 515 | 520 | 525 |
| Leu | Leu | Val | Tyr | Gly | Pro | Arg | Glu | Gly | Gly | Arg | Gly | Ala | Pro | Asp | | 530 | 535 | 540 |

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Pro | Phe | Leu | Gly | Val | Lys | Ala | Ala | Ala | Ala | Glu | Leu | Glu | Arg | Arg | |
| | | | | 545 | | | | | | 550 | | | | 555 | |
| Tyr | Pro | Gly | Thr | Arg | Leu | Ala | Trp | Leu | Ala | Val | Arg | Ala | Glu | Ala | |
| | | | | 560 | | | | | 565 | | | | | 570 | |
| Pro | Ser | Gln | Val | Arg | Leu | Met | Asp | Val | Val | Ser | Lys | Lys | His | Pro | |
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| Val | Asp | Thr | Leu | Phe | Phe | Leu | Thr | Thr | Val | Trp | Thr | Arg | Pro | Gly | |
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| Pro | Glu | Val | Leu | Asn | Arg | Cys | Arg | Met | Asn | Ala | Ile | Ser | Gly | Trp | |
| | | | | 605 | | | | | 610 | | | | | 615 | |
| Gln | Ala | Phe | Phe | Pro | Val | His | Phe | Gln | Glu | Phe | Asn | Pro | Ala | Leu | |
| | | | | 620 | | | | | 625 | | | | | 630 | |
| Ser | Pro | Gln | Arg | Ser | Pro | Pro | Gly | Pro | Pro | Gly | Ala | Gly | Pro | Asp | |
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| Pro | Pro | Ser | Pro | Pro | Gly | Ala | Asp | Pro | Ser | Arg | Gly | Ala | Pro | Ile | |
| | | | | 650 | | | | | 655 | | | | | 660 | |
| Gly | Gly | Arg | Phe | Asp | Arg | Gln | Ala | Ser | Ala | Glu | Gly | Cys | Phe | Tyr | |
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| Asn | Ala | Asp | Tyr | Leu | Ala | Ala | Arg | Ala | Arg | Leu | Ala | Gly | Glu | Leu | |
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| Ala | Gly | Gln | Glu | Glu | Glu | Glu | Ala | Leu | Glu | Gly | Leu | Glu | Val | Met | |
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| Asp | Val | Phe | Leu | Arg | Phe | Ser | Gly | Leu | His | Leu | Phe | Arg | Ala | Val | |
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| Glu | Pro | Gly | Leu | Val | Gln | Lys | Phe | Ser | Leu | Arg | Asp | Cys | Ser | Pro | |
| | | | | 725 | | | | | 730 | | | | | 735 | |
| Arg | Leu | Ser | Glu | Glu | Leu | Tyr | His | Arg | Cys | Arg | Leu | Ser | Asn | Leu | |
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| Glu | Gly | Leu | Gly | Gly | Arg | Ala | Gln | Leu | Ala | Met | Ala | Leu | Phe | Glu | |
| | | | | 755 | | | | | 760 | | | | | 765 | |
| Gln | Glu | Gln | Ala | Asn | Ser | Thr | | | | | | | | | |
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<211> 318

<212> PRT

<213> Homo Sapien

<400> 341

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| 1 | | | | 5 | | | | | 10 | | | | | 15 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Ile | Phe | Cys | Ala | Leu | Ile | Thr | Met | Leu | Gly | His | Ile | Arg | Ile |
| | | | | 20 | | | | | 25 | | | | | 30 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | His | Gly | Asn | Arg | Met | His | His | His | Glu | His | His | His | Leu | Gln |
| | | | | 35 | | | | | 40 | | | | | 45 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Pro | Asn | Lys | Glu | Asp | Ile | Leu | Lys | Ile | Ser | Glu | Asp | Glu | Arg |
| | | | | 50 | | | | | 55 | | | | | 60 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Glu | Leu | Ser | Lys | Ser | Phe | Arg | Val | Tyr | Cys | Ile | Ile | Leu | Val |
| | | | | 65 | | | | | 70 | | | | | 75 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Lys | Pro | Lys | Asp | Val | Ser | Leu | Trp | Ala | Ala | Val | Lys | Glu | Thr | Trp |
| | | | | 80 | | | | | 85 | | | | | 90 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Thr | Lys | His | Cys | Asp | Lys | Ala | Glu | Phe | Phe | Ser | Ser | Glu | Asn | Val |
| | | | | 95 | | | | | 100 | | | | | 105 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Lys | Val | Phe | Glu | Ser | Ile | Asn | Met | Asp | Thr | Asn | Asp | Met | Trp | Leu |
| | | | | 110 | | | | | 115 | | | | | 120 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Met | Arg | Lys | Ala | Tyr | Lys | Tyr | Ala | Phe | Asp | Lys | Tyr | Arg | Asp |
| | | | | 125 | | | | | 130 | | | | | 135 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gln | Tyr | Asn | Trp | Phe | Phe | Leu | Ala | Arg | Pro | Thr | Thr | Phe | Ala | Ile |
| | | | | 140 | | | | | 145 | | | | | 150 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Glu | Asn | Leu | Lys | Tyr | Phe | Leu | Leu | Lys | Lys | Asp | Pro | Ser | Gln |
| | | | | 155 | | | | | 160 | | | | | 165 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Phe | Tyr | Leu | Gly | His | Thr | Ile | Lys | Ser | Gly | Asp | Leu | Glu | Tyr |
| | | | | 170 | | | | | 175 | | | | | 180 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Gly | Met | Glu | Gly | Gly | Ile | Val | Leu | Ser | Val | Glu | Ser | Met | Lys |
| | | | | 185 | | | | | 190 | | | | | 195 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Arg | Leu | Asn | Ser | Leu | Leu | Asn | Ile | Pro | Glu | Lys | Cys | Pro | Glu | Gln |
| | | | | 200 | | | | | 205 | | | | | 210 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Gly | Met | Ile | Trp | Lys | Ile | Ser | Glu | Asp | Lys | Gln | Leu | Ala | Val |
| | | | | 215 | | | | | 220 | | | | | 225 |

Cys Leu Lys Tyr Ala Gly Val Phe Ala Glu Asn Ala Glu Asp Ala

| | 230 | 235 | 240 |
|-----------------|---------------------|---------------------|-----|
| Asp Gly Lys Asp | Val Phe Asn Thr Lys | Ser Val Gly Leu Ser | Ile |
| | 245 | 250 | 255 |
| Lys Glu Ala Met | Thr Tyr His Pro Asn | Gln Val Val Glu Gly | Cys |
| | 260 | 265 | 270 |
| Cys Ser Asp Met | Ala Val Thr Phe Asn | Gly Leu Thr Pro Asn | Gln |
| | 275 | 280 | 285 |
| Met His Val Met | Met Tyr Gly Val Tyr | Arg Leu Arg Ala Phe | Gly |
| | 290 | 295 | 300 |
| His Ile Phe Asn | Asp Ala Leu Val Phe | Leu Pro Pro Asn Gly | Ser |
| | 305 | 310 | 315 |

Asp Asn Asp

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ctggttcttc cttgcacg 18

<210> 344

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<210> 345

<211> 50

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<400> 347
ccctcatgta ccggctcc 18

<210> 348
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<210> 349
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<210> 351

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<210> 352

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<210> 353

<211> 48

<212> DNA

<213> Artificial Sequence

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<210> 355

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<212> DNA
<213> Homo Sapien

<400> 376

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tgctttctctt cccaaatgtt cttatggact gttgctggga tccccatcct 200
attttctcagt gcctgtttca tcaccagatg tgttgtgaca tttcgcatct 250
ttcaaacctg tgatgagaaa agttttcagc tacctgagaa tttcacagag 300
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gaactgggaa tattttcaat ccagctgcta cttcttttct actgacacca 400
tttctctgggc gttaagttta aagaactgct cagccatggg ggctcacctg 450
gtgggttatca actcacagga ggagcaggaa ttcctttcct acaagaaacc 500
taaaatgaga gagtttttta ttggactgtc agaccagggtt gtcgagggtc 550
agtggcaatg ggtggacggc acacctttga caaagtctct gagcttctgg 600
gatgtagggg agcccaacaa catagctacc ctggaggact gtgccaccat 650
gagagactct tcaaacccaa ggcaaaattg gaatgatgta acctgtttcc 700
tcaattattht tcggatttgt gaaatggtag gaataaatcc tttgaacaaa 750
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agagcaagaa catggccaca cccaccgccc cacacgagaa atttgtgcgc 850
tgaacttcaa aggacttcat aagtatttgt tactctgata caaataaaaa 900
taagtagttt taaatgttaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 950
aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaa 997
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<210> 377

<211> 219

<212> PRT

<213> Homo Sapien

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Cys Phe Ser Ser Gln Met Phe Leu Trp Thr Val Ala Gly Ile Pro
                20                      25                      30

Ile Leu Phe Leu Ser Ala Cys Phe Ile Thr Arg Cys Val Val Thr
                35                      40                      45
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| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| Phe | Arg | Ile | Phe | Gln | Thr | Cys | Asp | Glu | Lys | Lys | Phe | Gln | Leu | Pro | |
| | | | | 50 | | | | | 55 | | | | | 60 | |
| Glu | Asn | Phe | Thr | Glu | Leu | Ser | Cys | Tyr | Asn | Tyr | Gly | Ser | Gly | Ser | |
| | | | | 65 | | | | | 70 | | | | | 75 | |
| Val | Lys | Asn | Cys | Cys | Pro | Leu | Asn | Trp | Glu | Tyr | Phe | Gln | Ser | Ser | |
| | | | | 80 | | | | | 85 | | | | | 90 | |
| Cys | Tyr | Phe | Phe | Ser | Thr | Asp | Thr | Ile | Ser | Trp | Ala | Leu | Ser | Leu | |
| | | | | 95 | | | | | 100 | | | | | 105 | |
| Lys | Asn | Cys | Ser | Ala | Met | Gly | Ala | His | Leu | Val | Val | Ile | Asn | Ser | |
| | | | | 110 | | | | | 115 | | | | | 120 | |
| Gln | Glu | Glu | Gln | Glu | Phe | Leu | Ser | Tyr | Lys | Lys | Pro | Lys | Met | Arg | |
| | | | | 125 | | | | | 130 | | | | | 135 | |
| Glu | Phe | Phe | Ile | Gly | Leu | Ser | Asp | Gln | Val | Val | Glu | Gly | Gln | Trp | |
| | | | | 140 | | | | | 145 | | | | | 150 | |
| Gln | Trp | Val | Asp | Gly | Thr | Pro | Leu | Thr | Lys | Ser | Leu | Ser | Phe | Trp | |
| | | | | 155 | | | | | 160 | | | | | 165 | |
| Asp | Val | Gly | Glu | Pro | Asn | Asn | Ile | Ala | Thr | Leu | Glu | Asp | Cys | Ala | |
| | | | | 170 | | | | | 175 | | | | | 180 | |
| Thr | Met | Arg | Asp | Ser | Ser | Asn | Pro | Arg | Gln | Asn | Trp | Asn | Asp | Val | |
| | | | | 185 | | | | | 190 | | | | | 195 | |
| Thr | Cys | Phe | Leu | Asn | Tyr | Phe | Arg | Ile | Cys | Glu | Met | Val | Gly | Ile | |
| | | | | 200 | | | | | 205 | | | | | 210 | |
| Asn | Pro | Leu | Asn | Lys | Gly | Lys | Ser | Leu | | | | | | | |
| | | | | 215 | | | | | | | | | | | |

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<400> 423

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| Met | Ala | Leu | Arg | Arg | Pro | Pro | Arg | Leu | Arg | Leu | Cys | Ala | Arg | Leu |
| 1 | | | | 5 | | | | | 10 | | | | | 15 |

| | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Pro | Asp | Phe | Phe | Leu | Leu | Leu | Leu | Phe | Arg | Gly | Cys | Leu | Ile | Gly |
| | | | | 20 | | | | | 25 | | | | | 30 |

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 Phe Glu Ser Val Glu Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr 50 55 60
 Ser Asp Pro Arg Ile Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr 65 70 75
 Thr Tyr Val Phe Phe Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly 80 85 90
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 Thr Arg Arg Asp Ser Ala Leu Tyr Arg Cys Glu Val Val Ala Arg 110 115 120
 Asn Asp Arg Lys Glu Ile Asp Glu Ile Val Ile Glu Leu Thr Val 125 130 135
 Gln Val Lys Pro Val Thr Pro Val Cys Arg Val Pro Lys Ala Val 140 145 150
 Pro Val Gly Lys Met Ala Thr Leu His Cys Gln Glu Ser Glu Gly 155 160 165
 His Pro Arg Pro His Tyr Ser Trp Tyr Arg Asn Asp Val Pro Leu 170 175 180
 Pro Thr Asp Ser Arg Ala Asn Pro Arg Phe Arg Asn Ser Ser Phe 185 190 195
 His Leu Asn Ser Glu Thr Gly Thr Leu Val Phe Thr Ala Val His 200 205 210
 Lys Asp Asp Ser Gly Gln Tyr Tyr Cys Ile Ala Ser Asn Asp Ala 215 220 225
 Gly Ser Ala Arg Cys Glu Glu Gln Glu Met Glu Val Tyr Asp Leu 230 235 240
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 Leu Ala Leu Ile Thr Leu Gly Ile Cys Cys Ala Tyr Arg Arg Gly 260 265 270
 Tyr Phe Ile Asn Asn Lys Gln Asp Gly Glu Ser Tyr Lys Asn Pro 275 280 285
 Gly Lys Pro Asp Gly Val Asn Tyr Ile Arg Thr Asp Glu Glu Gly 290 295 300
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